transporter 1, ATP-binding cassette, sub-family B (MDR/TAP)	BF976260	2.23
RAD51 homolog C (S. cerevisiae)	AF029669	2.23
NADH dehydrogenase (ubiquinone) Fe-S protein 7 (20kD) (NADH-coenzyme Q reductase)	BC005954	2.23
ribosomal protein S8	NM_001012	2.23
vitamin A responsive; cytoskeleton related	N92494	2.23
spectrin SH3 domain binding protein 1	BF673013	2.23
ubiquinol-cytochrome c reductase, Rieske iron-sulfur polypeptide 1	BC000649	2.23
non-canonical ubiquitin conjugating enzyme 1	AF151039	2.23
aldo-keto reductase family 7, member A3 (aflatoxin aldehyde reductase)	NM_012067	2.23
peptide deformylase-like protein	NM_022341	2.23
nuclear cap binding protein subunit 2, 20kD	NM_007362	2.23
KIAA0225 protein	AW206115	2.22
amplified in osteosarcoma	AI683900	2.22
hypothetical protein FLJ10547	AL049795	2.22
putative translation initiation factor	BF246436	2.22
anaphase-promoting complex subunit 2	NM_013366	2.22
transmembrane protein 4	NM_014255	2.22
ES1 (zebrafish) protein, human homolog of	NM_004649	2.22
cell division cycle 27	NM_001256	2.22
KIAA0759 protein	AB018302	2.22
myosin IXB	AF143684	2.22
KIAA0673 protein	AB014573	2.22
X-prolyl aminopeptidase (aminopeptidase P) 2, membrane-bound	AL023653	2.22
ornithine decarboxylase 1	NM_002539	2.22
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ARP1 actin-related protein 1 homolog B, centractin beta (yeast)	NM_005735	2.22
tubulin, alpha 1 (testis specific)	K03460	2.22
KIAA0304 gene product	NM_014727	2.22
KIAA0542 gene product	AB011114	2.22
cyclin B2	NM_004701	2.21
MAX protein	AA723514	2.21
protein phosphatase 2 (formerly 2A), regulatory subunit B (PR 52), alpha isoform	NM_002717	2.21
DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 8 (RNA helicase)	NM_004941	2.21
sialyltransferase 7D ((alpha-N-acetylneuraminyl-2,3-beta-galactosyl-1,3)-N-acetyl galactosaminide alpha-2,6-sialyltransferase)	AB035172	2.21
heterogeneous nuclear ribonucleoprotein A3	NM_005758	2.21
CCAAT/enhancer binding protein (C/EBP), gamma	NM_001806	2.21
amino-terminal enhancer of split	NM_001130	2.21
CGI-152 protein	NM_020410	2.21
mitochondrial ribosomal protein L24	NM_024540	2.21
acetyl-Coenzyme A transporter	NM_004733	2.21
ATPase, H+ transporting, lysosomal (vacuolar proton pump) 9kD	AI862255	2.20
platelet/endothelial cell adhesion molecule (CD31 antigen)	M37780	2.20
pyruvate dehydrogenase (lipoamide) alpha 1	NM_000284	2.20
talin 1	NM_006289	2.20
actin, beta	NM_001101	2.20
Consensus includes gb:Z95126 /DEF=Human DNA sequence from PAC 30P20 on chromosome Xq21.1-Xq21.3. Contains set pseudogene, ESTs and STS /FEA=CDS /DB_XREF=gi:2342581 /U	Z95126	2.20
prion protein interacting protein	BC001072	2.20
hypothetical protein FLJ11937	NM_022765	2.19
hypothetical protein DKFZp434E1723	AI199589	2.19
ribosomal protein S4, X-linked	NM_001007	2.19
Consensus includes gb:AU144066 /FEA=EST /DB_XREF=gi:11005587 /DB_XREF=est:AU144066 /CLONE=HEMBA1000798 /UG=Hs.285519 Homo sapiens cDNA FLJ11904 fis, clone HEMBB	AU144066	2.19
hypothetical protein FLJ11021 similar to splicing factor, arginine/serine-rich 4	NM_023012	2.19
hypothetical protein	BF038366	2.19
cytochrome c oxidase subunit VIb pseudogene-3	AL031594	2.19
v-Ha-ras Harvey rat sarcoma viral oncogene homolog	NM_005343	2.19
RNA polymerase II transcriptional regulation mediator (Med6, S. cerevisiae, homolog of)	AF074723	2.19
polymerase (RNA) II (DNA directed) polypeptide C (33kD)	AC004382	2.19
CGI-36 protein	NM_015963	2.18
DKFZP586A011 protein	NM_015416	2.18
DKFZP586G011 protein	BG289527	2.18
mitofusin 1	U95822	2.18
hypothetical protein MGC3731	NM_024313	2.18
chromosome 11 hypothetical protein ORF4	NM_020155	2.18

adaptor-related protein complex 2, alpha 2 subunit	BC006155	2.18
SET translocation (myeloid leukemia-associated)	M93651	2.18
tumor differentially expressed 1	U49188	2.18
eukaryotic translation elongation factor 2	AJ004246	2.17
hypothetical protein FLJ10377	NM_018077	2.17
3-hydroxy-3-methylglutaryl-Coenzyme A synthase 1 (soluble)	NM_002130	2.17
phosphatidylinositol 4-kinase, catalytic, beta polypeptide	NM_002651	2.17
proteasome (prosome, macropain) subunit, alpha type 6	BC002979	2.17
postmeiotic segregation increased 2-like 8	NM_005394	2.17
ubiquinol-cytochrome c reductase (6.4kD) subunit	AC005943	2.17
ligase I, DNA, ATP-dependent	NM_000234	2.17
pre-B-cell colony-enhancing factor	BF575514	2.17
sperm associated antigen 9	AB011088	2.17
glycyl-tRNA synthetase	D30658	2.17
DnaJ (Hsp40) homolog, subfamily B, member 12	NM_017626	2.17
general transcription factor II, i	BC004472	2.17
GM2 ganglioside activator protein	AL513583	2.17
guanine nucleotide binding protein (G protein), beta polypeptide 1	AF070603	2.16
autoantigen	NM_014329	2.16
RNA binding motif protein 8A	NM_005105	2.16
putative translation initiation factor	AL537707	2.16
eukaryotic translation initiation factor 3, subunit 5 (epsilon, 47kD)	NM_003754	2.16
hypothetical protein MGC11279	NM_024326	2.16
aldo-keto reductase family 7, member A2 (aflatoxin aldehyde reductase)	NM_003689	2.16
chloride intracellular channel 4	AF109196	2.16
annexin A11	NM_001157	2.16
ATPase, H+ transporting, lysosomal (vacuolar proton pump) 16kD	M62762	2.16
phosphoprotein regulated by mitogenic pathways	NM_025195	2.16
glutathione synthetase	NM_000178	2.16
RNA-binding protein regulatory subunit	NM_007262	2.16
protein phosphatase 2A 48 kDa regulatory subunit	NM_013239	2.16
conserved helix-loop-helix ubiquitous kinase	AF080157	2.16
nuclear transcription factor, X-box binding 1	AW291398	2.16
coatomer protein complex, subunit alpha	U24105	2.15
sorbitol dehydrogenase	L29008	2.15
eukaryotic translation elongation factor 1 alpha 1	AL515273	2.15
adaptor-related protein complex 3, delta 1 subunit	AF130042	2.15
metallothionein 2A	NM_005953	2.15
adenylosuccinate synthase	AA628948	2.15
hypothetical protein MGC2718	NM_024067	2.15
KIAA0867 protein	NM_014938	2.15
hypothetical protein FLJ20312	NM_017761	2.15
microtubule-associated protein, RP/EB family, member 1	NM_012325	2.14
Ran GTPase activating protein 1	BE379408	2.14
CD164 antigen, sialomucin	AF263279	2.14
CGI-128 protein	NM_016062	2.14
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postmeiotic segregation increased 2-like 9	NM_005395	2.14
RPB5-mediating protein	AB006572	2.13
phosphorylase kinase, beta	BG149218	2.13
cytochrome c oxidase subunit VIII	NM_004074	2.13
GPAA1P anchor attachment protein 1 homolog (yeast)	NM_003801	2.13
hypothetical protein MGC2749	BC006479	2.13
alpha-actinin-2-associated LIM protein	AF002280	2.13
membrane metallo-endopeptidase (neutral endopeptidase, enkephalinase, CALLA, CD10)	NM_007287	2.13
F-box only protein 9	AJ954041	2.13
CGI-125 protein	NM_016060	2.13
hypothetical protein FLJ10631	NM_018161	2.13
signal peptidase 12kDa	NM_014041	2.13
general transcription factor IIIC, polypeptide 5 (63kD)	NM_012087	2.13
seven in absentia homolog 2 (Drosophila)	U76248	2.13
heterogeneous nuclear ribonucleoprotein C (C1/C2)	AA664258	2.13
integrin-linked kinase	NM_004517	2.13

X-ray repair complementing defective repair in Chinese hamster cells 5 (double-strand-break rejoining; Ku autoantigen, 80kD)	J04977	2.13
dynactin 4	AI038068	2.13
cyclin B1	BE407516	2.13
deoxythymidylate kinase (thymidylate kinase)	NM_012145	2.13
Sec23 homolog A (S. cerevisiae)	NM_006364	2.13
ubinuclein 1	NM_016936	2.12
keratin, hair, acidic, 3A	NM_004138	2.12
actin, beta	AA809056	2.12
adducin 1 (alpha)	BE898639	2.12
prostate cancer overexpressed gene 1	NM_003627	2.12
amyloid beta (A4) precursor-like protein 2	BC004371	2.12
RecQ protein-like 4	NM_004260	2.12
CGI-51 protein	NM_015380	2.12
CTP synthase	NM_001905	2.12
glutamyl-prolyl-tRNA synthetase	NM_004446	2.12
signal sequence receptor, alpha (translocon-associated protein alpha)	AI016620	2.11
RAB7, member RAS oncogene family	BG261416	2.11
polypyrimidine tract binding protein (heterogeneous nuclear ribonucleoprotein I)	NM_002819	2.11
sialyltransferase 9 (CMP-NeuAc:lactosylceramide alpha-2,3-sialyltransferase; GM3 synthase)	NM_003896	2.11
KIAA0164 gene product	BE963370	2.11
BRCA1 associated protein	AF035620	2.11
McKusick-Kaufman syndrome	NM_018848	2.11
polypyrimidine tract binding protein (heterogeneous nuclear ribonucleoprotein I)	BC004383	2.11
guanine nucleotide binding protein (G protein), beta 5	AF300650	2.11
adenomatosis polyposis coli	AI375486	2.10
hypothetical protein, clone 24751	AF070530	2.10
hypothetical protein FLJ10975	NM_018288	2.10
DKFZP586J0619 protein	BF513430	2.10
zinc finger protein 22 (KOX 15)	AA744771	2.10
NOL1R protein	AI284845	2.10
microtubule-associated protein 4	NM_002375	2.10
acid phosphatase 1, soluble	NM_004300	2.10
dolichyl-phosphate mannosyltransferase polypeptide 3	NM_018973	2.09
glucose regulated protein, 58kD	D83485	2.09
DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 3	R60068	2.09
pyruvate dehydrogenase (lipoamide) beta	M34055	2.09
prefoldin 2	NM_012394	2.09
proline synthetase co-transcribed (bacterial homolog)	AL136616	2.09
neighbor of COX4	NM_006067	2.09
molybdenum cofactor synthesis 2	NM_004531	2.09
EH-domain containing 1	AF001434	2.09
hypothetical protein FLJ21324	NM_021941	2.09
phosphofructokinase, liver	NM_002626	2.09
spectrin, alpha, non-erythrocytic 1 (alpha-fodrin)	U83867	2.09
gb:AF092132.1 /DEF=Homo sapiens PAK2 mRNA, complete cds. /FEA=mRNA /PROD=PAK2 /DB_XREF=gi:5138913 /UG=Hs.284275 Homo sapiens PAK2 mRNA, complete cds /FL=gb:AF09:	AF092132	2.09
calcium binding protein Cab45 precursor	NM_016547	2.09
growth factor, augmenter of liver regeneration (ERV1 homolog, S. cerevisiae)	AF124604	2.09
Rho-specific guanine nucleotide exchange factor p114	AB011093	2.08
zinc finger protein 289, IDI regulated	BE299671	2.08
up-regulated by BCG-CWS	AL049963	2.08
hypothetical protein FLJ12542	NM_024899	2.08
KIAA0872 protein	AI629033	2.08
low molecular mass ubiquinone-binding protein (9.5kD)	NM_014402	2.08
transcription factor Dp-1	R60866	2.08
CREBBP/EP300 inhibitory protein 1	AF109873	2.08
guanine nucleotide binding protein (G protein), beta 5	NM_006578	2.07
hypothetical protein	R51077	2.07
hypothetical protein FLJ20758	NM_017952	2.07
DKFZP434J154 protein	NM_016003	2.07
phosphoribosylglycinamide formyltransferase, phosphoribosylglycinamide synthetase, phosphoribosylaminoimidazole synthetase	NM_000819	2.07
gb:BC001233.1 /DEF=Homo sapiens, Similar to KIAA0092 gene product, clone MGC:4896, mRNA, complete cds. /FEA=mRNA /PROD=Similar to KIAA0092 gene product /DB_XREF=gi:12654	BC001233	2.07
suppression of tumorigenicity 7	NM_018412	2.07
hypothetical protein FLJ20607	NM_017899	2.07

interleukin enhancer binding factor 3, 90kD	BG032366	2.06
basic leucine-zipper protein BZAP45	AL518328	2.06
membrane interacting protein of RGS16	NM_016641	2.06
gb:Z25425.1 /DEF=H.sapiens protein-serinethreonine kinase gene, complete CDS. /FEA=mRNA /PROD=protein-serinethreonine kinase /DB_XREF=gi:405728 /FL=gb:Z25425.1	Z25425	2.06
suppression of tumorigenicity 13 (colon carcinoma) (Hsp70 interacting protein)	BE866412	2.06
POP4 (processing of precursor , S. cerevisiae) homolog	NM_006627	2.06
intercellular adhesion molecule 2	AA126728	2.06
jumping translocation breakpoint	AF151056	2.06
microtubule-associated protein like echinoderm EMAP	AI582285	2.06
Consensus includes gb:BF037823 /FEA=EST /DB_XREF=gi:10746209 /DB_XREF=est:601461994F1 /CLONE=IMAGE:3865487 /UG=Hs.9683 DnaJ (Hsp40) homolog, subfamily C, member 3	BF037823	2.06
polyamine-modulated factor 1	NM_007221	2.06
progestin induced protein	AF006010	2.06
tumor differentially expressed 1	BC006088	2.06
SET translocation (myeloid leukemia-associated)	AI278616	2.06
proteasome (prosome, macropain) 26S subunit, non-ATPase, 13	NM_002817	2.06
sorting nexin 1	NM_003099	2.06
RABSA, member RAS oncogene family	NM_004162	2.06
basic leucine zipper transcription factor, ATF-like	AC007182	2.06
pituitary tumor-transforming 1 interacting protein	NM_004339	2.06
sialidase 1 (lysosomal sialidase)	U84246	2.06
postmeiotic segregation increased 2-like 8	AB017004	2.05
A kinase (PRKA) anchor protein 1	U34074	2.05
promyelocytic leukemia	AF230401	2.05
chromosome 21 open reading frame 4	NM_006134	2.05
tubulin, alpha 1 (testis specific)	AL565074	2.05
alkylglycerone phosphate synthase	NM_003659	2.05
FUS interacting protein (serine-arginine rich) 2	AU130523	2.05
hypothetical protein MGC2491	NM_024040	2.05
lysyl-tRNA synthetase	AF285758	2.05
Clp1-interacting zinc finger protein	AA054734	2.05
protein tyrosine phosphatase, non-receptor type 22 (lymphoid)	NM_015967	2.05
oxoglutarate (alpha-ketoglutarate) dehydrogenase (lipoamide)	NM_002541	2.05
sperm associated antigen 7	NM_004890	2.05
actin, beta	X00351	2.04
CD2 antigen (cytoplasmic tail) binding protein 2	NM_006110	2.04
agrin	AI424797	2.04
serine/threonine kinase 38	BE048506	2.04
Consensus includes gb:AI653730 /FEA=EST /DB_XREF=gi:4737709 /DB_XREF=est:wb36f12.x1 /CLONE=IMAGE:2307791 /UG=Hs.279009 matrix Gla protein	AI653730	2.04
hyaluronan-mediated motility receptor (RHAMM)	U29343	2.04
KIAA0999 protein	NM_025164	2.04
heat shock factor binding protein 1	NM_001537	2.04
antigen identified by monoclonal antibody Ki-67	AU152107	2.04
sulfotransferase family, cytosolic, 1C, member 2	NM_006588	2.04
KIAA0184 protein	D80006	2.04
hypothetical protein FLJ11183	NM_018351	2.04
superoxide dismutase 1, soluble (amyotrophic lateral sclerosis 1 (adult))	NM_000454	2.03
MIL1 protein	NM_015367	2.03
HLA class II region expressed gene KE2	NM_014260	2.03
inhibitor of growth family, member 1-like	AI186701	2.03
stearoyl-CoA desaturase (delta-9-desaturase)	AA678241	2.03
phosphomevalonate kinase	NM_006556	2.03
gb:BC000265.1 /DEF=Homo sapiens, clone MGC:3182, mRNA, complete cds. /FEA=mRNA /PROD=Unknown (protein for MGC:3182) /DB_XREF=gi:12653008 /UG=Hs.315417 Homo sapiens	BC000265	2.03
tumor necrosis factor receptor superfamily, member 6b, decoy	AL121845	2.03
hypothetical protein FLJ11200	NM_018359	2.03
prosaposin (variant Gaucher disease and variant metachromatic leukodystrophy)	NM_002778	2.03
FK506 binding protein 5	NM_004117	2.03
EH-domain containing 1	NM_006795	2.03
postmeiotic segregation increased 2-like 1	D38435	2.03
N-myc downstream-regulated gene 3	BE300252	2.03
ash2 (absent, small, or homeotic)-like (Drosophila)	AB020982	2.03
Mad4 homolog	AA831438	2.02
ATP-binding cassette, sub-family E (OABP), member 1	AI002002	2.02
PTD016 protein	NM_016125	2.02

KIAA1118 protein	AB029041	2.02
mitochondrial ribosomal protein L42	BE782148	2.02
hypothetical protein FLJ10036	NM_017975	2.02
phosphotidylinositol transfer protein	NM_006224	2.02
calcium-regulated heat-stable protein (24kD)	NM_014316	2.02
HSPC038 protein	NM_016096	2.02
E2IG2 protein	NM_016565	2.02
polymerase (DNA directed), delta 1, catalytic subunit (125kD)	NM_002691	2.02
hypothetical protein	NM_015932	2.02
CD74 antigen (invariant polypeptide of major histocompatibility complex, class II antigen-associated)	X01144	2.02
ADP-ribosyltransferase (NAD+; poly (ADP-ribose) polymerase)-like 1	AL359763	2.02
Consensus includes gb:AL109942 /DEF=Human DNA sequence from clone RP3-473J16 on chromosome 6q25.3-26 Contains 3 end of the MAP3K4 (mitogen-activated protein kinase kinase ki	AL109942	2.02
actin, gamma 1	AL515810	2.02
methionine adenosyltransferase II, alpha	NM_005911	2.02
ribosomal protein S17	NM_001021	2.01
RNA binding motif protein 5	NM_005778	2.01
hypothetical protein PP2447	BG399562	2.01
protein translocation complex beta	NM_006808	2.01
v-src sarcoma (Schmidt-Ruppin A-2) viral oncogene homolog (avian)	NM_005417	2.01
hypothetical protein FLJ22104	AK000684	2.01
acetyl-Coenzyme A acetyltransferase 1 (acetoacetyl Coenzyme A thiolase)	NM_000019	2.01
glycerol-3-phosphate dehydrogenase 2 (mitochondrial)	U36310	2.01
guanidinoacetate N-methyltransferase	NM_000156	2.01
tumor necrosis factor (ligand) superfamily, member 8	NM_001244	2.01
v-abl Abelson murine leukemia viral oncogene homolog 1	NM_005157	2.01
polypyrimidine tract binding protein (heterogeneous nuclear ribonucleoprotein I)	BC002397	2.01
phosphodiesterase 4A, cAMP-specific (phosphodiesterase E2 dunce homolog, Drosophila)	NM_006202	2.01
OKFZP4348168 protein	AK001363	2.01
Consensus includes gb:AK002076.1 /DEF=Homo sapiens cDNA FLJ11214 fis, clone PLACE1007990. /FEA=mRNA /DB_XREF=gi:7023738 /UG=Hs.281616 Homo sapiens cDNA FLJ11214 fis,	AK002076	2.01
CD53 antigen	NM_000560	2.01
talfalin-interacting protein	AL080147	2.01
PTD009 protein	NM_016146	2.01
ADP-ribosyltransferase (NAD+; poly (ADP-ribose) polymerase)	M32721	2.01
mitochondrial ribosomal protein S18A	NM_018135	2.01
similar to Bos taurus P14 protein	NM_030877	2.01
zinc finger protein 195	AI476267	2.01
homocysteine-inducible, endoplasmic reticulum stress-inducible, ubiquitin-like domain member 1	AF217990	2.01
embryonic ectoderm development	AF080227	2.01
mannose phosphate isomerase	NM_002435	2.00
DAZ associated protein 1	NM_018959	2.00
coilin	NM_004645	2.00
high-mobility group (nonhistone chromosomal) protein 17-like 3	NM_006353	2.00
DAZ associated protein 2	AL534321	2.00
granulin	AK023348	2.00
cell membrane glycoprotein, 110000M(r) (surface antigen)	NM_007002	2.00
aspartylglucosaminidase	W27131	2.00
gene from NF2/meningioma region of 22q12	BC003615	2.00
ribosomal protein S10	AA320764	2.00
KIAA1393 protein	AB037814	2.00
golgi associated, gamma adaptin ear containing, ARF binding protein 2	BE646414	2.00
HCR (a-helix coiled-coil rod homologue)	AI588986	1.99
breast cancer 1, early onset	AF005068	1.99
enoyl Coenzyme A hydratase, short chain, 1, mitochondrial	NM_004092	1.99
KIAA0971 protein	AK021557	1.99
B-cell translocation gene 1, anti-proliferative	NM_001731	1.99
hypothetical protein FLJ13057 similar to germ cell-less	NM_022471	1.99
KIAA0040 gene product	T79953	1.99
proteasome (prosome, macropain) activator subunit 2 (PA28 beta)	NM_002818	1.99
RAN binding protein 7	AA939270	1.99
prostatic binding protein	BE969671	1.99
hematopoietic protein 1	BC001604	1.99
interleukin-1 receptor-associated kinase 1	NM_001569	1.99
casein kinase 2, alpha 1 polypeptide	AI161318	1.99

short-chain dehydrogenase/reductase 1	NM_004753	1.99
Consensus includes gb:AF052117.1 /DEF=Homo sapiens clone 23809 mRNA sequence. /FEA=mRNA /DB_XREF=gi:3360424 /UG=Hs.6932 Homo sapiens clone 23809 mRNA sequence	AF052117	1.99
Wilms' tumour 1-associating protein	BC000383	1.99
RNA binding motif protein 8A	AI738479	1.99
ataxia telangiectasia mutated (includes complementation groups A, C and D)	NM_000051	1.99
KIAA0365 gene product	AV738039	1.99
DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 17 (72kD)	AW188131	1.99
metastasis associated 1	BC006177	1.99
hypothetical protein FLJ13984	NM_024770	1.99
60S ribosomal protein L30 isolog	NM_016304	1.98
similar to S. cerevisiae Sec6p and R. norvegicus rsec6	AF055006	1.98
epilepsy, progressive myoclonus type 2, Lafora disease (laforin)	BC005286	1.98
hypothetical protein	NM_019048	1.98
UDP-glucose pyrophosphorylase 2	NM_006759	1.98
phosphodiesterase 6D, cGMP-specific, rod, delta	AI001626	1.98
Consensus includes gb:M21388.1 /DEF=Human unproductively rearranged Ig mu-chain mRNA V-region (VD), 5 end, clone mu-3A1A. /FEA=mRNA /GEN=IGHM /DB_XREF=gi:185160 /UG=H	M21388	1.98
hypothetical protein	AI081779	1.98
NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 5 (13kD, B13)	NM_005000	1.98
putative nucleolar RNA helicase	NM_019082	1.98
heterogeneous nuclear ribonucleoprotein K	AI701949	1.98
tumor protein D52-like 2	NM_003288	1.98
H2A histone family, member E	NM_021066	1.98
SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily a, member 4	AK026573	1.98
ribosomal protein S3A	AI925635	1.98
uridine monophosphate kinase	BC002906	1.98
dishevelled, dsh homolog 1 (Drosophila)	AF006011	1.98
synaptic nuclei expressed gene 2	NM_015180	1.97
postmeiotic segregation increased 2-like 2	AA161026	1.97
ras-related C3 botulinum toxin substrate 1 (rho family, small GTP binding protein Rac1)	BG292367	1.97
uroporphyrinogen III synthase (congenital erythropoietic porphyria)	NM_000375	1.97
zinc finger protein 266	AA868898	1.97
RAD17 homolog (S. pombe)	AF085736	1.97
vacuolar protein sorting 26 (yeast)	NM_004896	1.97
polycystic kidney disease 1 (autosomal dominant)	AK023376	1.97
uncharacterized hematopoietic stem/progenitor cells protein MDS026	AI138993	1.96
survival of motor neuron protein interacting protein 1	AB037701	1.96
RecQ protein-like (DNA helicase Q1-like)	NM_002907	1.96
eukaryotic translation initiation factor 3, subunit 1 (alpha, 35kD)	AL031313	1.96
Consensus includes gb:N95443 /FEA=EST /DB_XREF=gi:1267753 /DB_XREF=est:zb81c12.s1 /CLONE=IMAGE:310006 /UG=Hs.19180 Homo sapiens mRNA; cDNA DKFZp564E122 (from clo	N95443	1.96
cyclin-dependent kinase inhibitor 2A (melanoma, p16, inhibits CDK4)	U38945	1.96
Nit protein 2	NM_020202	1.96
CGI-65 protein	NM_016013	1.96
karyopherin alpha 4 (importin alpha 3)	U93240	1.96
hypothetical protein MGC20727	AI879381	1.96
small nuclear ribonucleoprotein D1 polypeptide (16kD)	BC001721	1.96
protein phosphatase 5, catalytic subunit	NM_006247	1.96
conserved gene amplified in osteosarcoma	NM_005730	1.96
similar to S. cerevisiae SSM4	BF000409	1.96
ribonuclease MI, large subunit	NM_006397	1.96
DKFZP566C084 protein	NM_015510	1.96
WD repeat domain 1	NM_017491	1.96
pM5 protein	N39536	1.96
non-metastatic cells 2, protein (NM23B) expressed in	NM_002512	1.96
hypothetical protein FLJ13449	NM_024546	1.96
protein kinase C substrate 80K-H	NM_002743	1.96
gb:AF293339.1 /DEF=Homo sapiens ornithine decarboxylase antizyme 4 mRNA, complete cds. /FEA=mRNA /PROD=ornithine decarboxylase antizyme 4 /DB_XREF=gi:11024341 /FL=gb:AF	AF293339	1.95
hypothetical protein FLJ21839	NM_021831	1.95
SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily a, member 4	D26156	1.95
CD47 antigen (Rh-related antigen, integrin-associated signal transducer)	Z25521	1.95
carbonyl reductase	NM_016286	1.95
RNA polymerase I subunit	AF008442	1.95
arsenate resistance protein ARS2	AI523895	1.95
Consensus includes gb:BE379006 /FEA=EST /DB_XREF=gi:9324371 /DB_XREF=est:601236272F1 /CLONE=IMAGE:3608680 /UG=Hs.99766 Homo sapiens mRNA; cDNA DKFZp564I0323 (h	BE379006	1.95

ribosomal protein S12	AI799007	1.95
Consensus includes gb:AI671747 /FEA=EST /DB_XREF=gi:4851478 /DB_XREF=est:wa05e06.x1 /CLONE=IMAGE:2297218 /UG=Hs.81360 Homo sapiens cDNA: FLJ21927 fis, clone HEP041	AI671747	1.95
acid phosphatase 1, soluble	BE872974	1.95
staufen, RNA binding protein (Drosophila)	BC000830	1.95
hypothetical protein FLJ10450	NM_018095	1.95
G protein beta subunit-like	NM_022372	1.95
BRCA1 associated RING domain 1	NM_000465	1.95
hypothetical protein FLJ20452	NM_017828	1.95
CD24 antigen (small cell lung carcinoma cluster 4 antigen)	L33930	1.95
neuronal protein 17.3	NM_019056	1.94
zinc finger protein 43 (HTF6)	NM_003423	1.94
ribosomal protein L4	AI953886	1.94
proteasome (prosome, macropain) subunit, alpha type, 1	NM_002786	1.94
calcium/calmodulin-dependent protein kinase IG	AW016039	1.94
proteasome (prosome, macropain) subunit, beta type, 2	NM_002794	1.94
putative membrane protein	AF277194	1.94
O-linked N-acetylglucosamine (GlcNAc) transferase (UDP-N-acetylglucosamine:polypeptide-N-acetylglucosaminyl transferase)	U77413	1.94
Rho-associated, coiled-coil containing protein kinase 1	AV683882	1.94
CGI-31 protein	NM_015959	1.94
integrin-linked kinase-associated serine/threonine phosphatase 2C	AY024365	1.94
Ope-interacting protein 2	AL050353	1.94
GABA(A) receptor-associated protein	NM_007278	1.94
jumping translocation breakpoint	NM_006694	1.94
ROD1 regulator of differentiation 1 (S. pombe)	AW190873	1.94
sema domain, immunoglobulin domain (Ig), transmembrane domain (TM) and short cytoplasmic domain, (semaphorin) 4D	NM_006378	1.94
cyclic AMP phosphoprotein, 19 kD	BC003418	1.94
RAP2B, member of RAS oncogene family	NM_002886	1.93
potassium intermediate/small conductance calcium-activated channel, subfamily N, member 3	NM_002249	1.93
ankyrin repeat and SOCS box-containing 1	AF055024	1.93
Cluster Incl. H59033:yr40e10.r1 Homo sapiens cDNA, 5 end /clone=IMAGE-207786 /clone_end=5 /gb=H59033 /gi=1011865 /ug=Hs.6700 /len=604	H59033	1.93
trophinin associated protein (tastin)	NM_005480	1.93
suppressor of Ty 4 homolog 1 (S. cerevisiae)	BC002802	1.93
ribosomal protein L36a	NM_001001	1.93
hypothetical protein FLJ10359	NM_018072	1.93
DHHC1 protein	BE542551	1.93
HSPC009 protein	NM_014019	1.93
KIAA0217 protein	AU154358	1.93
phosphoglycerate mutase 1 (brain)	NM_002629	1.93
protein tyrosine phosphatase, receptor type, O	U20489	1.93
matrin 3	AA577695	1.93
core-binding factor, beta subunit	AF294326	1.93
KIAA0456 protein	AA742261	1.93
Consensus includes gb:AK025862.1 /DEF=Homo sapiens cDNA: FLJ22209 fis, clone HRC01496. /FEA=mRNA /DB_XREF=gi:10438505 /UG=Hs.288528 Homo sapiens cDNA: FLJ22209 fis, cl	AK025862	1.93
BUB3 budding uninhibited by benzimidazoles 3 homolog (yeast)	AU160695	1.93
hypothetical protein FLJ10315	NM_018056	1.93
FK506 binding protein 4 (59kD)	AA894574	1.92
death-associated protein	NM_004394	1.92
hypothetical protein FLJ20850	NM_017967	1.92
Consensus includes gb:AK023154.1 /DEF=Homo sapiens cDNA FLJ13092 fis, clone NT2RP3002147. /FEA=mRNA /DB_XREF=gi:10434948 /UG=Hs.172035 hypothetical protein similar to m	AK023154	1.92
eukaryotic translation elongation factor 1 alpha 1	BE786672	1.92
alpha integrin binding protein 63	NM_007216	1.92
gb:AF061735.1 /DEF=Homo sapiens clone 023e08 My032 protein mRNA, complete cds. /FEA=mRNA /PROD=My032 protein /DB_XREF=gi:12002005 /UG=Hs.307099 Homo sapiens clone	AF061735	1.92
ORF	NM_015853	1.92
KIAA0170 gene product	AI673553	1.92
HSPC144 protein	NM_014174	1.92
hypothetical protein DKFZp56400523	AL527334	1.92
cyclin I	AF135162	1.92
cofilin 1 (non-muscle)	NM_005507	1.92
endosulfine alpha	BC000436	1.92
squamous cell carcinoma antigen recognised by T cells 3	AW173076	1.92
phosphoprotein enriched in astrocytes 15	BC002426	1.92
growth hormone 2	AF006060	1.92
chromosome 1 open reading frame 16	NM_014837	1.92

clone FLB4739	AA133341	1.92
putative translation initiation factor	AL523310	1.92
ATP-binding cassette, sub-family F (GCN20), member 1	NM_001090	1.92
diphtheria toxin receptor (heparin-binding epidermal growth factor-like growth factor)	NM_001945	1.92
UDP-Gal:betaGlcNAc beta 1,4- galactosyltransferase, polypeptide 6	AF097159	1.91
hypothetical protein DKFZp762M115	BC002477	1.91
translocase of inner mitochondrial membrane 17 homolog A (yeast)	BC004439	1.91
proteasome (prosome, macropain) subunit, alpha type, 5	NM_002790	1.91
ribosomal protein S27 (metallopanstimulin 1)	NM_001030	1.91
gamma tubulin ring complex protein (76p gene)	BC000966	1.91
steroid sulfatase (microsomal), arylsulfatase C, isozyme S	AI122754	1.91
chromatin assembly factor 1, subunit A (p150)	BF062223	1.91
C3H-type zinc finger protein; similar to D. melanogaster muscleblind B protein	NM_005757	1.91
angio-associated, migratory cell protein	NM_001087	1.91
adaptor-related protein complex 2, alpha 2 subunit	AA877641	1.91
DNA segment on chromosome X (unique) 9928 expressed sequence	NM_004699	1.91
hypothetical protein	NM_016407	1.91
hypothetical protein	BE314601	1.91
Consensus includes gb:AU144378 /FEA=EST /DB_XREF=gi:11005899 /DB_XREF=est:AU144378 /CLONE=HEMBA1001734 /UG=Hs.296650 Homo sapiens cDNA FLJ11475 fis, clone HEMBA	AU144378	1.91
polypyrimidine tract binding protein (heterogeneous nuclear ribonucleoprotein I)	BF690062	1.91
isoprenylcysteine carboxyl methyltransferase	NM_012405	1.91
tyrosine 3-monooxygenase/tryptophan 5-monooxygenase activation protein, zeta polypeptide	NM_003406	1.91
polymerase (RNA) II (DNA directed) polypeptide E (25kD)	NM_002695	1.91
FKSO6 binding protein 4 (59kD)	NM_002014	1.90
Consensus includes gb:AW304232 /FEA=EST /DB_XREF=gi:6713921 /DB_XREF=est:zv82g01.x1 /CLONE=IMAGE:2825040 /UG=Hs.181357 laminin receptor 1 (67kD, ribosomal protein SA)	AW304232	1.90
protein phosphatase 3 (formerly 2B), regulatory subunit B (19kD), alpha isoform (calcineurin B, type I)	NM_000945	1.90
Mdm2, transformed 3T3 cell double minute 2, p53 binding protein (mouse)	AI276888	1.90
EH-domain containing 1	AL579035	1.90
cysteinyl-tRNA synthetase	AI769685	1.90
basic leucine-zipper protein BZAP45	NM_014670	1.90
hypothetical protein MGC14258	U79256	1.90
Consensus includes gb:AL121975 /DEF=Human DNA sequence from clone RP3-422B11 on chromosome 6p11.2-12.3 Contains ESTs, STSs and GSSs. Contains part of the PRIM2A gene for tw	AL121975	1.90
putative receptor protein	NM_003876	1.90
PAP-1 binding protein	NM_031288	1.90
myosin regulatory light chain interacting protein	NM_013262	1.90
high-mobility group (nonhistone chromosomal) protein isoforms I and Y	NM_002131	1.90
serine hydroxymethyltransferase 1 (soluble)	L23928	1.90
phosphatidylserine synthase 1	NM_014754	1.90
hypothetical protein FLJ10297	NM_018049	1.90
IQ motif containing GTPase activating protein 1	D29640	1.90
neurocalcin delta	AF251061	1.90
secretory carrier membrane protein 2	NM_005697	1.90
ubiquinol-cytochrome c reductase core protein II	AV727381	1.90
glutathione-S-transferase like; glutathione transferase omega	NM_004832	1.90
transportin-SR	AF145029	1.89
gb:NM_000970.1 /DEF=Homo sapiens ribosomal protein L6 (RPL6), mRNA. /FEA=mRNA /GEN=RPL6 /PROD=ribosomal protein L6 /DB_XREF=gi:4506656 /UG=Hs.174131 ribosomal prote	NM_000970	1.89
HSPC156 protein	NM_014178	1.89
low density lipoprotein receptor defect C complementing	NM_007357	1.89
requiem, apoptosis response zinc finger gene	NM_006268	1.89
cyclin G2	NM_004354	1.89
DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 9 (RNA helicase A, nuclear DNA helicase II; leukophysin)	BE561014	1.89
RecQ protein-like (DNA helicase Q1-like)	BC001052	1.89
plasminogen activator, urokinase receptor	AC005525	1.89
HBS1-like (S. cerevisiae)	BC001465	1.89
transcription factor-like 4	BF056105	1.89
neuroepithelial cell transforming gene 1	NM_005863	1.89
BRCA1-interacting protein 1	AF360549	1.89
PRNC, apoptosis, WT1, regulator	NM_002583	1.89
hypothetical protein FLJ13962	NM_024862	1.89
chromosome 20 open reading frame 80	AL441988	1.89
hypothetical protein FLJ10099	NM_017994	1.89
inositol polyphosphate-5-phosphatase, 75kD	AI039084	1.89
C9orf10 protein	BE963765	1.88

ELK1, member of ETS oncogene family	M25269	1.88
hypothetical protein DKFZp434G0522	NM_017566	1.88
G protein-coupled receptor 30	U63917	1.88
transcription factor A, mitochondrial	NM_012251	1.88
palmitoyl-protein thioesterase 1 (ceroid-lipofuscinosis, neuronal 1, infantile)	NM_000310	1.88
GM2 ganglioside activator protein	X62078	1.88
ATP citrate lyase	NM_001096	1.88
myeloid cell leukemia sequence 1 (BCL2-related)	NM_021960	1.88
likely ortholog of mouse tubulin alpha 4	NM_025019	1.88
thymidine kinase 1, soluble	NM_003258	1.88
KIAA0522 protein	BF571239	1.88
proteasome (prosome, macropain) 26S subunit, ATPase, 4	NM_006503	1.88
general transcription factor IIH, polypeptide 1 (62kD subunit)	NM_005316	1.88
NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 10 (42kD)	NM_004544	1.88
non-canonical ubiquitin conjugating enzyme 1	NM_016021	1.88
hypothetical protein FLJ23221	NM_024579	1.88
protein-L-isoaspartate (D-aspartate) O-methyltransferase	M93008	1.88
DnaJ (Hsp40) homolog, subfamily A, member 2	AF011793	1.88
KIAA0121 gene product	D50911	1.87
DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 17 (72kD)	U59321	1.87
ubiquitin B	NM_018955	1.87
high-mobility group (nonhistone chromosomal) protein 17	BC003689	1.87
hypothetical protein	NM_019005	1.87
RNA polymerase I subunit	NM_004875	1.87
cyclin A2	NM_001237	1.87
neighbor of A-kinase anchoring protein 95	NM_014371	1.87
N-acetylglucosamine kinase	NM_017567	1.87
exosome component Rrp41	AI571298	1.87
hypothetical protein from EUROIMAGE 1987170	AL136935	1.87
Rab geranylgeranyltransferase, alpha subunit	NM_004581	1.87
cullin 4B	AI650819	1.87
Era G-protein-like 1 (E. coli)	AL562733	1.87
dUTP pyrophosphatase	U90223	1.87
granulin	NM_002087	1.87
ubiquitin specific protease 10	BC000263	1.87
steroid sulfatase (microsomal), arylsulfatase C, isozyme S	J04964	1.87
SMC2 structural maintenance of chromosomes 2-like 1 (yeast)	NM_006444	1.87
ATPase, aminophospholipid transporter (APLT), Class I, type 8A, member 1	AB013452	1.87
ribosomal protein S17	BC004886	1.87
DKFZP434D156 protein	NM_015530	1.87
hypothetical protein R33729_1	AL524093	1.87
DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 11 (CHL1-like helicase homolog, S. cerevisiae)	U33833	1.87
hypothetical protein FLJ22301	NM_024836	1.86
PRO1575 protein	NM_014092	1.86
argininosuccinate lyase	NM_000048	1.86
transcription factor A, mitochondrial	BE552470	1.86
ubiquitin-conjugating enzyme E2G 2 (UBC7 homolog, yeast)	BC001738	1.86
ubiquitin carboxyl-terminal esterase L3 (ubiquitin thiolesterase)	NM_006002	1.86
chromodomain protein, Y chromosome-like	NM_004824	1.86
E1A binding protein p300	AI459462	1.86
hypothetical protein FLJ21986	NM_024913	1.86
amino acid transporter 2	NM_018976	1.86
zinc finger protein, subfamily 1A, 1 (Ikaros)	BG540504	1.86
gb:AF116705.1 /DEF=Homo sapiens PRO2489 mRNA, complete cds. /FEA=mRNA /PROD=PRO2489 /DB_XREF=gi:7959908 /FL=gb:AF116705.1	AF116705	1.86
hypothetical protein RG083M05.2	NM_021167	1.86
desmoglein 1	NM_001942	1.86
actin, gamma 1	AW190090	1.86
KIAA0153 protein	BF965437	1.86
hypothetical protein FLJ11021 similar to splicing factor, arginine/serine-rich 4	BE396879	1.86
hypothetical protein	BF038366	1.85
high-mobility group (nonhistone chromosomal) protein 1	BE311760	1.85
erbb2 interacting protein	NM_018695	1.85
suppressor of Ty 5 homolog (S. cerevisiae)	NM_003169	1.85

Sec3-like	AK023461	1.85
RAB3 GTPase-ACTIVATING PROTEIN	AI040009	1.85
vesicle docking protein p115	NM_003715	1.85
glucosamine-6-phosphate isomerase	NM_005471	1.85
beta-amyloid binding protein precursor	AF353990	1.85
Xq28, 2000bp sequence contg. ORF	NM_017518	1.85
solute carrier family 21 (organic anion transporter), member 8	NM_019844	1.85
ubiquitin-conjugating enzyme E2L 3	AI819709	1.85
cytochrome c oxidase subunit IV isoform 1	AA854966	1.85
sperm associated antigen 4	NM_003116	1.85
cell death regulator aven	NM_020371	1.85
protein tyrosine phosphatase, receptor type, f polypeptide (PTPRF), interacting protein (liprin), alpha 1	BG033593	1.85
delta sleep inducing peptide, immunoreactor	AL110191	1.85
erythrocyte membrane protein band 7.2 (stomatin)	AF130103	1.85
cytochrome c oxidase subunit VIIa polypeptide 2 (liver)	AC004544	1.85
reticulon 2	AF004222	1.85
outer dense fibre of sperm tails 2	AF053970	1.85
protease, serine, 25	NM_013247	1.84
DKFZP586E1519 protein	AK022874	1.84
H2A histone family, member Z	NM_002106	1.84
squamous cell carcinoma antigen recognised by T cells	NM_005146	1.84
MCM4 minichromosome maintenance deficient 4 (S. cerevisiae)	AA604621	1.84
heterogeneous nuclear ribonucleoprotein A3	AA528233	1.84
poly(A) binding protein, cytoplasmic 1	AI734929	1.84
HSPC162 protein	NM_014183	1.84
ataxia telangiectasia and Rad3 related	U49844	1.84
nuclear factor (erythroid-derived 2)-like 2	NM_006164	1.84
RSP1-like protein	AA887480	1.84
MAGEF1 protein	NM_022149	1.84
Consensus includes gb:AW873564 /FEA=EST /DB_XREF=gi:8007617 /DB_XREF=est:ho62c10.x1 /CLONE=IMAGE:3041970 /UG=Hs.155485 huntingtin interacting protein 2	AW873564	1.84
Consensus includes gb:AW129021 /FEA=EST /DB_XREF=gi:6116965 /DB_XREF=est:xe93h03.x1 /CLONE=IMAGE:2615477 /UG=Hs.243537 ESTs, Highly similar to AC005061 1 similar to A/	AW129021	1.84
Consensus includes gb:BC004409.1 /DEF=Homo sapiens, clone IMAGE:3638994, mRNA, partial cds. /FEA=mRNA /PROD=Unknown (protein for IMAGE:3638994) /DB_XREF=gi:13325181 /L	BC004409	1.84
hypothetical protein PP591	AL562282	1.84
protein tyrosine phosphatase, receptor type, f polypeptide (PTPRF), interacting protein (liprin), alpha 1	U22815	1.84
nuclear factor of kappa light polypeptide gene enhancer in B-cells inhibitor, beta	AI684894	1.84
CGI-146 protein	AF229834	1.84
hypothetical protein KIAA1164	AU121431	1.84
ubiquitin-conjugating enzyme HBUCE1	W74577	1.84
proteasome (prosome, macropain) subunit, alpha type, 1	BC005932	1.84
transcription factor 3 (E2A immunoglobulin enhancer binding factors E12/E47)	AA768906	1.84
cell division cycle 2-like 1 (PITSLRE proteins)	AI767436	1.84
ADP-ribosylation factor-like 1	NM_001177	1.84
Rac/Cdc42 guanine exchange factor (GEF) 6	D25304	1.83
syntaxin 7	AI566082	1.83
peptidylprolyl isomerase F (cyclophilin F)	BC005020	1.83
glutaredoxin (thioltransferase)	AF162769	1.83
KIAA0540 protein	AB011112	1.83
telomeric repeat binding factor 2	NM_005652	1.83
KIAA0293 protein	AB006631	1.83
KIAA0573 protein	BC005374	1.83
hypothetical protein FLJ20783	NM_017958	1.83
elaC homolog 2 (E. coli)	NM_018127	1.83
hypothetical protein	NM_016404	1.83
hypothetical protein 38408_6	NM_014551	1.83
Clustar Incl. AA628140:af97e06.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-1055746 /clone_end=3 /gb=AA628140 /gi=2540527 /ug=Hs.126089 /len=302	AA628140	1.83
ring finger protein 1	AL031228	1.83
ribosomal protein L14	AA838274	1.83
gb:U72518.1 /DEF=Human destrin-2 pseudogene mRNA, complete cds. /FEA=mRNA /PROD=destrin-2 /DB_XREF=gi:1673523 /UG=Hs.199299 Human destrin-2 pseudogene mRNA, compl	U72518	1.83
proteasome (prosome, macropain) activator subunit 3 (PA28 gamma; Ki)	BC001423	1.83
COP9 (constitutive photomorphogenic, Arabidopsis, homolog) subunit 7A	AF193844	1.83
apoptosis-associated speck-like protein containing a CARD	BC004470	1.83
hypothetical protein FLJ10747	NM_018202	1.83
nuclear distribution gene C (A.nidulans) homolog	AF241788	1.83

scratch homolog 1, zinc finger protein (Drosophila)	NM_031309	1.83
N-myristoyltransferase 1	AF020500	1.82
eukaryotic translation initiation factor 1A	BG149698	1.82
CGI-48 protein	NM_016001	1.82
NS1-binding protein	NM_006469	1.82
hypothetical protein FLJ13114	H27225	1.82
gb:BC002746.1 /DEF=Homo sapiens, Similar to dodecenoyl-Coenzyme A delta isomerase (3,2 trans-enoyl-Coenzyme A isomerase), clone MGC:3903, mRNA, complete cds. /FEA=mRNA /PR(	BC002746	1.82
DKFZP564D116 protein	BE962354	1.82
mutS homolog 3 (E. coli)	NM_002439	1.82
hypothetical protein MGC11308	BF690150	1.82
DKFZP586B0519 protein	NM_015456	1.82
myosin, light polypeptide, regulatory, non-sarcomeric (20kD)	NM_006471	1.82
diacylglycerol O-acyltransferase homolog 1 (mouse)	NM_012079	1.82
protein phosphatase 2A, regulatory subunit B' (PR 53)	NM_021131	1.82
origin recognition complex, subunit 5-like (yeast)	NM_002553	1.82
poly(rC) binding protein 2	NM_005016	1.82
tRNA-guanine transglycosylase	NM_031209	1.82
butyrate-induced transcript 1	NM_016395	1.82
cytochrome c oxidase subunit Vb	AI557312	1.82
DKFZp566O133 protein	AL050050	1.82
Consensus includes gb:AK024450.1 /DEF=Homo sapiens mRNA for FLJ00042 protein, partial cds. /FEA=mRNA /GEN=FLJ00042 /PROD=FLJ00042 protein /DB_XREF=gi:10440413 /UG=Hs.	AK024450	1.82
high-mobility group (nonhistone chromosomal) protein 17-like 3	BC001282	1.82
mitochondrial ribosomal protein L34	AB049652	1.82
ornithine decarboxylase antizyme 1	D87914	1.82
ribosomal protein L13	AI186735	1.82
KIAA0737 gene product	AL117508	1.82
proteasome (prosome, macropain) activator subunit 1 (PA28 alpha)	NM_006263	1.82
golgi SNAP receptor complex member 1	NM_004871	1.82
phosphatidylserine receptor	AK021780	1.82
dihydrofolate reductase	BC003584	1.82
gene with multiple splice variants near HD locus on 4p16.3	NM_003704	1.82
protein phosphatase 3 (formerly 2B), catalytic subunit, beta isoform (calcineurin A beta)	M29550	1.82
chromosome 14 open reading frame 1	AL136658	1.82
tyrosylprotein sulfotransferase 2	NM_003595	1.82
hypothetical protein FLJ13912	AI421812	1.82
KIAA0092 gene product	AI123527	1.81
mitogen-activated protein kinase kinase 2	NM_030662	1.81
MAD, mothers against decapentaplegic homolog 2 (Drosophila)	NM_005901	1.81
ectonucleotide pyrophosphatase/phosphodiesterase 4 (putative function)	AW194947	1.81
leptin receptor overlapping transcript-like 1	AF161461	1.81
L-3-hydroxyacyl-Coenzyme A dehydrogenase, short chain	AF001903	1.81
hypothetical protein DKFZp762A227	AI630178	1.81
KIAA0947 protein	BC004902	1.81
hypothetical protein FLJ10326	NM_018060	1.81
succinate-CoA ligase, GDP-forming, beta subunit	BF593940	1.81
ribosomal protein S6	BC000524	1.81
proteasome (prosome, macropain) 26S subunit, non-ATPase, 11	BF432873	1.81
bone marrow stromal cell antigen 1	NM_004334	1.81
ras homolog gene family, member G (rho G)	NM_001665	1.81
hypothetical protein FLJ22570	NM_024872	1.81
copper chaperone for superoxide dismutase	NM_005125	1.81
TAF12 RNA polymerase II, TATA box binding protein (TBP)-associated factor, 20 kD	D50544	1.81
gb:Z25430.1 /DEF=H.sapiens protein-serinethreonine kinase gene, complete CDS. /FEA=mRNA /PROD=protein-serinethreonine kinase /DB_XREF=gi:405738 /FL=gb:Z25430.1	Z25430	1.81
hypothetical protein MGC3062	NM_024065	1.81
DnaJ (Hsp40) homolog, subfamily C, member 7	NM_003315	1.81
HIF-1 responsive RTP801	NM_019058	1.80
eukaryotic translation initiation factor 2B, subunit 5 (epsilon, 82kD)	U23028	1.80
KIAA0462 protein	AB007931	1.80
mitogen-activated protein kinase phosphatase x	NM_020185	1.80
hypothetical protein	NM_015383	1.80
zinc finger protein 258	NM_007167	1.80
protein phosphatase 2, regulatory subunit B (B56), delta isoform	AB000635	1.80
Rho-associated, coiled-coil containing protein kinase 2	AL049383	1.80

zinc finger protein 259, pseudogene	Z95118	1.80
hypothetical protein FLJ20452	BE313890	1.80
integrin, beta 1 (fibronectin receptor, beta polypeptide, antigen CD29 includes MDF2, MSK12)	AA215854	1.80
SWAP-70 protein	AB014540	1.80
v-rel reticuloendotheliosis viral oncogene homolog (avian)	NM_002908	1.80
cullin 4B	NM_003588	1.80
vacuolar protein sorting 16 (yeast)	NM_022575	1.79
breast cancer 2, early onset	NM_000059	1.79
eukaryotic translation initiation factor 2, subunit 1 (alpha, 35kD)	BC002513	1.79
MAD, mothers against decapentaplegic homolog 2 (Drosophila)	U65019	1.79
hypothetical protein FLJ22175	AA551370	1.79
DKFZP586D0623 protein	BC000758	1.79
CHK1 checkpoint homolog (S. pombe)	NM_001274	1.79
CHK2 checkpoint homolog (S. pombe)	BC004207	1.79
hypothetical protein FLJ21080	NM_022743	1.79
ubiquitin specific protease 1	NM_003368	1.79
neuroblastoma RAS viral (v-ras) oncogene homolog	NM_002524	1.79
UDP-GlcNAc:betaGal beta-1,3-N-acetylglucosaminyltransferase 1	NM_006577	1.79
ubiquitin carboxyl-terminal esterase L1 (ubiquitin thiolesterase)	NM_004181	1.79
complement component 1, q subcomponent binding protein	AU151801	1.79
ATPase, H+ transporting, lysosomal (vacuolar proton pump) membrane sector associated protein M8-9	NM_005765	1.79
polymerase (RNA) III (DNA directed) (62kD)	U93867	1.79
proteasome (prosome, macropain) 26S subunit, non-ATPase, 7 (Mov34 homolog)	NM_002811	1.79
phorbol-12-myristate-13-acetate-induced protein 1	AI857639	1.79
GCN5 general control of amino-acid synthesis 5-like 2 (yeast)	NM_021078	1.79
ATP synthase, H+ transporting, mitochondrial F0 complex, subunit c (subunit 9) isoform 3	NM_001689	1.79
HeLa cyclin-dependent kinase 2 interacting protein	NM_016550	1.79
cytochrome c oxidase subunit VIIb pseudogene 1	AF042164	1.78
thyroid hormone receptor interactor 8	AI694023	1.78
similar to Caenorhabditis elegans protein C42C1.9	AL568449	1.78
zinc finger protein 95 homolog (mouse)	BF196931	1.78
KIAA0251 protein	AK025504	1.78
ring finger protein 14	NM_004290	1.78
Sec23 homolog B (S. cerevisiae)	NM_006363	1.78
ADP-ribosylation factor 3	NM_001659	1.78
serine/arginine repetitive matrix 2	NM_016333	1.78
DKFZp564I157 protein	NM_018457	1.78
clone HQ0477 PRO0477p	NM_016360	1.78
ATPase, H+ transporting, lysosomal (vacuolar proton pump), member J	BC003564	1.78
sirtuin silent mating type information regulation 2 homolog 7 (S. cerevisiae)	NM_016538	1.78
malic enzyme 2, NAD(+)-dependent, mitochondrial	BC000147	1.78
baculoviral IAP repeat-containing 5 (survivin)	NM_001168	1.78
KIAA1085 protein	AL136943	1.78
delta sleep inducing peptide, immunoreactor	NM_004089	1.78
hypothetical protein DKFZp564K142 similar to implantation-associated protein	AL136636	1.77
c6.1A	S72931	1.77
transcriptional regulator interacting with the PHS-bromodomain 2	BG107456	1.77
protein phosphatase 1, regulatory (inhibitor) subunit 16B	AB020630	1.77
somatostatin receptor 5	NM_001053	1.77
aldo-keto reductase family 7, member A2 (aflatoxin aldehyde reductase)	AI144075	1.77
transmembrane 9 superfamily member 1	BE899402	1.77
KIAA0874 protein	AW572909	1.77
diazepam binding inhibitor (GABA receptor modulator, acyl-Coenzyme A binding protein)	M15887	1.77
H326	AK026481	1.77
myogenic factor 3	BC000353	1.77
protein disulfide isomerase-related protein	BC001312	1.77
eukaryotic translation initiation factor 2B, subunit 1 (alpha, 26kD)	NM_001414	1.77
Consensus includes gb:N92498 /FEA=EST /DB_XREF=gi:1264807 /DB_XREF=est:zb28a04.s1 /CLONE=IMAGE:304878 /UG=Hs.326248 Homo sapiens cDNA: FLJ22071 fis, clone HEP11691	N92498	1.77
cystathionine-beta-synthase	BE613178	1.77
S100 calcium binding protein A4 (calcium protein, calvasculin, metastasin, murine placental homolog)	NM_002961	1.77
NY-REN-24 antigen	AI200896	1.77
ES1 (zebrafish) protein, human homolog of	D86062	1.77
mitochondrial ribosomal protein S28	NM_014018	1.77

transcription elongation factor B (SIII), polypeptide 1 (15kD, elongin C)	N89607	1.76
basic transcription factor 3, like 2	M90355	1.76
ubiquitin specific protease 13 (isopeptidase T-3)	NM_003940	1.76
pleckstrin homology, Sec7 and coiled/coil domains 2 (cytohesin-2)	BC004361	1.76
hypothetical protein FLJ22427	AK026080	1.76
splicing factor, arginine/serine-rich 2	BE866585	1.76
immediate early protein	NM_004907	1.76
myosin IE	NM_004998	1.76
a disintegrin and metalloproteinase domain 9 (meltrin gamma)	NM_003816	1.76
hypothetical protein FLJ22573	NM_024660	1.76
hypothetical protein similar to actin related protein 2/3 complex, subunit 5	NM_030978	1.76
hydroxyacyl glutathione hydrolase	NM_005326	1.76
DiGeorge syndrome critical region gene 2	AU150824	1.76
eukaryotic translation elongation factor 1 beta 2	NM_001959	1.76
RAN binding protein 3	AI689052	1.76
zinc finger protein 282	AW130128	1.76
hypothetical protein FLJ21908	NM_024604	1.76
SNARE protein	BE384482	1.76
citrate synthase	BC000105	1.76
replication factor C (activator 1) 4 (37kD)	NM_002916	1.76
ribosomal protein L17	NM_000985	1.76
ribonuclease P (38kD)	NM_006414	1.76
trinucleotide repeat containing 12	BE880591	1.76
aldolase C, fructose-bisphosphate	NM_005165	1.76
hypothetical protein MGC3133	NM_031287	1.76
phosphatidylinositol 4-kinase type II	AL561930	1.76
KIAA0317 gene product	AC007956	1.76
N-ethylmaleimide-sensitive factor	NM_006178	1.76
hexosaminidase A (alpha polypeptide)	AL523158	1.75
ADP-ribosylation factor-like 2	NM_001667	1.75
glutathione S-transferase M1	NM_000561	1.75
KIAA1033 protein	AK001657	1.75
solute carrier family 16 (monocarboxylic acid transporters), member 1	NM_003051	1.75
ADP-ribosylation factor 5	NM_001662	1.75
guanine nucleotide binding protein (G protein), alpha inhibiting activity polypeptide 3	J03198	1.75
polymerase (DNA directed), epsilon	AL080203	1.75
fatty-acid-Coenzyme A ligase, long-chain 1	NM_001995	1.75
bromodomain-containing 7	AI885109	1.75
myelin transcription factor 1	AB028973	1.75
serine/threonine kinase 17b (apoptosis-inducing)	NM_004226	1.75
hypothetical protein FLJ23282	AA907940	1.75
nuclear cap binding protein subunit 1, 80kD	BC001450	1.75
phosphorylase, glycogen; brain	NM_002862	1.75
dihydrolipoamide S-succinyltransferase (E2 component of 2-oxo-glutarate complex)	S72422	1.75
hypothetical nuclear factor SBBI22	NM_020395	1.75
ribosomal protein L12	NM_000976	1.75
tumor rejection antigen (gp96) 1	AI582238	1.75
mevalonate (diphospho) decarboxylase	AI189359	1.75
CGI-45 protein	NM_015999	1.75
KIAA0595 protein	NM_015062	1.75
hypothetical protein FLJ20721	NM_017941	1.74
matrin 3	NM_018834	1.74
PEST-containing nuclear protein	NM_020357	1.74
replication protein A1 (70kD)	BG398414	1.74
inositol polyphosphate phosphatase-like 1	NM_001567	1.74
hypothetical protein FLJ10581	NM_018146	1.74
chromosome 5 open reading frame 8	NM_006816	1.74
SM-11044 binding protein	NM_020123	1.74
propionyl Coenzyme A carboxylase, beta polypeptide	NM_000532	1.74
anaphase-promoting complex 1; meiotic checkpoint regulator	NM_022662	1.74
MCM3 minichromosome maintenance deficient 3 (S. cerevisiae) associated protein	AJ010089	1.74
TP53 target gene 1	AB007458	1.74
actin, gamma 1	BC001920	1.74

hypothetical protein FLJ12492	NM_024586	1.74
hepatitis B virus x-interacting protein (9.6kD)	NM_006402	1.74
nuclear factor (erythroid-derived 2)-like 1	AI361227	1.74
Golgi apparatus protein 1	AK025457	1.74
Consensus includes gb:AL034550 /DEF=Human DNA sequence from clone RP5-1184F4 on chromosome 20q11.1-11.23. Contains the 3 end of gene KIAA0978, two genes for novel proteins	AL034550	1.74
cyclin G associated kinase	NM_005255	1.74
zinc finger protein 9 (a cellular retroviral nucleic acid binding protein)	NM_003418	1.74
GLI-Kruppel family member HKR3	NM_005341	1.74
hypothetical protein FLJ12910	NM_024573	1.74
low density lipoprotein receptor-related protein 8, apolipoprotein e receptor	NM_017522	1.74
troponin C, slow	AF020769	1.74
similar to MRJ gene for a member of the DNAJ protein family (H. sapiens)	BC002446	1.74
EAP30 subunit of ELL complex	NM_007241	1.74
hypothetical protein FLJ12529	NM_024811	1.74
CD19 antigen	NM_001770	1.74
RAD17 homolog (S. pombe)	AF098533	1.74
Consensus includes gb:AK026977.1 /DEF=Homo sapiens cDNA: FLJ23324 fis, clone HEP12482, highly similar to HUMMYOHCB Human nonmuscle myosin heavy chain-B (MYH10) mRNA. /FE	AK026977	1.74
heat shock factor binding protein 1	AK026575	1.74
tubulin, alpha, ubiquitous	NM_006082	1.74
hypothetical protein FLJ22530	AI422099	1.74
pre-mRNA splicing factor 17	NM_015891	1.74
membrane metallo-endopeptidase (neutral endopeptidase, enkephalinase, CALLA, CD10)	AI433463	1.74
G-rich RNA sequence binding factor 1	BF034561	1.74
cell division cycle 34	BG476661	1.74
casein kinase 1, alpha 1	NM_001892	1.73
uncharacterized bone marrow protein BM042	NM_018458	1.73
nuclear receptor coactivator 3	U80737	1.73
inositol polyphosphate-5-phosphatase, 145kD	NM_005541	1.73
TAR (HIV) RNA binding protein 2	NM_004178	1.73
HSPC171 protein	BC003080	1.73
tetratricopeptide repeat domain 3	AU131711	1.73
proteasome (prosome, macropain) 26S subunit, ATPase, 5	AF035309	1.73
hypothetical protein FLJ20707	NM_017936	1.73
major histocompatibility complex, class II, DM beta	NM_002118	1.73
dendritic cell protein	NM_006360	1.73
BRCA1 associated protein-1 (ubiquitin carboxy-terminal hydrolase)	NM_004656	1.73
KIAA0244 protein	AI949220	1.73
survival of motor neuron protein interacting protein 1	NM_003616	1.73
stress 70 protein chaperone, microsome-associated, 60kD	NM_006948	1.73
dihydrolipoamide S-acetyltransferase (E2 component of pyruvate dehydrogenase complex)	BF978872	1.73
PI-3-kinase-related kinase SMG-1	BG256504	1.73
KIAA0247 gene product	NM_014734	1.73
chromosome 11 open reading frame 24	AA065185	1.73
tubulin-specific chaperone e	NM_003193	1.73
nucleolar protein family A, member 2 (H/ACA small nucleolar RNPs)	BC000009	1.73
small nuclear ribonucleoprotein 70kD polypeptide (RNP antigen)	NM_003089	1.73
granulin	BC000324	1.73
B-cell receptor-associated protein BAP29	NM_018844	1.73
casein kinase 1, gamma 3	NM_004384	1.73
succinate dehydrogenase complex, subunit C, integral membrane protein, 15kD	AF080579	1.72
SMART/HDAC1 associated repressor protein	NM_015001	1.72
COP9 homolog	BC003090	1.72
actin, gamma 1	NM_001614	1.72
KIAA0943 protein	AL080168	1.72
CCCTC-binding factor (zinc finger protein)	NM_006565	1.72
solute carrier family 6 (neurotransmitter transporter, GABA), member 13	NM_016615	1.72
bromodomain-containing 4	BF718610	1.72
tetratricopeptide repeat domain 1	NM_003314	1.72
histidine triad nucleotide binding protein	NM_005340	1.72
uroporphyrinogen decarboxylase	M14016	1.72
Consensus includes gb:BC003542.1 /DEF=Homo sapiens, clone IMAGE:3611719, mRNA, partial cds. /FEA=mRNA /PROD=Unknown (protein for IMAGE:3611719) /DB_XREF=gi:13097656 /L	BC003542	1.72
KIAA0746 protein	AB018289	1.72
dolichyl-P-Glc:Man9GlcNAc2-PP-dolichylglucosyltransferase	NM_013339	1.72

mitofusin 2	NM_014874	1.72
glutaryl-Coenzyme A dehydrogenase	NM_000159	1.72
ATPase, Ca++ transporting, cardiac muscle, slow twitch 2	M23114	1.72
MAD, mothers against decapentaplegic homolog 4 (Drosophila)	NM_005359	1.72
bromodomain adjacent to zinc finger domain, 1B	NM_023005	1.72
X-ray repair complementing defective repair in Chinese hamster cells 1	NM_006297	1.72
ferredoxin 1	M18003	1.72
proteasome (prosome, macropain) subunit, beta type, 6	BC000835	1.72
ATPase, H+ transporting lysosomal (vacuolar proton pump), member M	AF100741	1.72
isocitrate dehydrogenase 3 (NAD+) beta	NM_006899	1.72
metallothionein 1H	NM_005951	1.72
ferredoxin 1	NM_004109	1.72
heterogeneous nuclear ribonucleoprotein C (C1/C2)	AV725195	1.72
tumor necrosis factor receptor superfamily, member 17	NM_001192	1.72
hypothetical protein MGC2722	NM_024102	1.71
RE1-silencing transcription factor	NM_005612	1.71
protoporphyrinogen oxidase	NM_000309	1.71
PC2 (positive cofactor 2, multiprotein complex) glutamine/Q-rich-associated protein	AK000003	1.71
Arg/Abl-interacting protein ArgBP2	NM_021069	1.71
glyceraldehyde-3-phosphate dehydrogenase	M33197	1.71
NADH dehydrogenase (ubiquinone) Fe-S protein 2 (49kD) (NADH-coenzyme Q reductase)	NM_004550	1.71
eNOS interacting protein	NM_015953	1.71
staufen, RNA binding protein (Drosophila)	NM_004602	1.71
ADP-ribosylation factor-like 1	BE890745	1.71
hypothetical protein FLJ20897	AI613383	1.71
CGI-06 protein	NM_015937	1.71
zinedin	NM_013403	1.71
small fragment nuclease	NM_015523	1.71
forkhead box O1A (rhabdomyosarcoma)	NM_002015	1.71
galactokinase 2	NM_002044	1.71
KIAA0603 gene product	NM_014832	1.71
DNA (cytosine-5-)-methyltransferase 1	NM_001379	1.71
syntrophin, beta 1 (dystrophin-associated protein A1, 59kD, basic component 1)	NM_021021	1.71
N-acetyltransferase, homolog of S. cerevisiae ARD1	NM_003491	1.71
KIAA0431 protein	NM_015251	1.71
tRNA isopentenylpyrophosphate transferase	BE964125	1.71
Consensus includes gb:AL121916 /DEF=Human DNA sequence from clone RP1-189G13 on chromosome 20. Contains an RPL7A (60S ribosomal protein L7A) (SURF3) pseudogene, part of ar	AL121916	1.71
protein disulfide isomerase-related protein	NM_005742	1.71
transformer-2 alpha (htra-2 alpha)	AW978896	1.71
KIAA0930 protein	AK025608	1.71
formin-binding protein 17	AB011126	1.71
hypthetical protein PRO2389	NM_025230	1.71
cysteine-rich protein 1 (intestinal)	NM_001311	1.70
novel putative protein similar to YIL091C yeast hypothetical 84 kD protein from SGA1-KTR7	N30910	1.70
hypothetical protein FLJ11583	AA398062	1.70
collagen, type VI, alpha 1	M20776	1.70
ribosomal protein L12	AK026491	1.70
tubulin, beta polypeptide	BF971587	1.70
Notch homolog 2 (Drosophila)	AU158495	1.70
gb:NM_001514.2 /DEF=Homo sapiens general transcription factor IIB (GTF2B), mRNA. /FEA=mRNA /GEN=GTF2B /PROD=general transcription factor IIB /DB_XREF=gi:13435384 /FL=gb:N	NM_001514	1.70
protein kinase, lysine deficient 1	AI445745	1.70
mitochondrial ribosomal protein S18B	AL050361	1.70
SET binding factor 1	U93181	1.70
hypothetical protein PRO2730	AL137654	1.70
major histocompatibility complex, class I, A	AI923492	1.70
sorting nexin 6	NM_021249	1.70
ZAP3 protein	AI952986	1.70
postmeiotic segregation increased 2-like 8	AI375694	1.70
serine/threonine kinase 18	NM_014264	1.70
malate dehydrogenase 2, NAD (mitochondrial)	BC001917	1.70
NOL1R2 protein	AK021688	1.70
filamin B, beta (actin binding protein 278)	M62994	1.70
membrane cofactor protein (CD46, trophoblast-lymphocyte cross-reactive antigen)	NM_002389	1.70

ribosomal protein L21	NM_000982	1.70
L-3-hydroxyacyl-Coenzyme A dehydrogenase, short chain	NM_005327	1.70
protein tyrosine phosphatase, non-receptor type 18 (brain-derived)	NM_014369	1.70
BCL2/adenovirus E1B 19kD interacting protein 1	U15172	1.70
hypothetical protein bK1048E9.5	NM_022081	1.70
MAP/microtubule affinity-regulating kinase 3	AI745639	1.70
ribosomal protein L35a	BE968801	1.70
integrin cytoplasmic domain-associated protein 1	NM_004763	1.70
hypothetical protein FLJ11126	NM_018332	1.69
nuclear transcription factor Y, gamma	NM_014223	1.69
hypothetical protein MGC5378	BC000927	1.69
protein phosphatase 3 (formerly 2B), catalytic subunit, gamma isoform (calcineurin A gamma)	NM_005605	1.69
KIAA0052 protein	D29641	1.69
insulin-degrading enzyme	N22903	1.69
CDC23 (cell division cycle 23, yeast, homolog)	NM_004661	1.69
Consensus includes gb:AU144887 /FEA=EST /DB_XREF=gi:11006408 /DB_XREF=est:AU144887 /CLONE=HEMBA1003278 /UG=Hs.301800 Homo sapiens cDNA FLJ11568 fis, clone HEMBA	AU144887	1.69
Consensus includes gb:BG168283 /FEA=EST /DB_XREF=gi:12674986 /DB_XREF=est:602340822F1 /CLONE=IMAGE:4448789 /UG=Hs.82202 ribosomal protein L17	BG168283	1.69
uroporphyrinogen decarboxylase	AW614435	1.69
Consensus includes gb:AL050136.1 /DEF=Homo sapiens mRNA; cDNA DKFZp586L141 (from clone DKFZp586L141). /FEA=mRNA /DB_XREF=gi:4884346 /UG=Hs.140945 Homo sapiens mR	AL050136	1.69
KIAA0544 protein	AB011116	1.69
hypothetical protein	BC000979	1.69
Consensus includes gb:AL121777 /DEF=Human DNA sequence from clone RP5-1057D4 on chromosome 20 Contains a Spermidine synthase (SPDSY) pseudogene, a CpG island, ESTs, STSs a	AL121777	1.69
serine palmitoyltransferase, long chain base subunit 2	NM_004863	1.69
disca, large (Drosophila) homolog 1	AW139131	1.69
ankyrin 1, erythrocytic	AF005213	1.69
cytochrome c oxidase subunit VIa polypeptide 1	NM_004373	1.69
dipeptidylpeptidase III	NM_005700	1.69
JM1 protein	NM_014008	1.69
differentially expressed in FDCP 6 homolog (mouse)	NM_022047	1.69
cleavage and polyadenylation specific factor 5, 25 kD subunit	NM_007006	1.69
chromosome 14 open reading frame 2	AF116639	1.69
KIAA0092 gene product	AA918224	1.69
NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 8 (19kD, ASHI)	NM_005004	1.69
TCF3 (E2A) fusion partner (in childhood Leukemia)	NM_013342	1.69
myotubularin related protein 4	AL042220	1.69
isoprenylcysteine carboxyl methyltransferase	AL578502	1.69
brefeldin A-inhibited guanine nucleotide-exchange protein 1	AK025637	1.69
melanoma-associated antigen recognised by cytotoxic T lymphocytes	NM_006428	1.68
DKFZp434J1813 protein	BG168666	1.68
Consensus includes gb:AI348009 /FEA=EST /DB_XREF=gi:4085215 /DB_XREF=est:qp61b09.x1 /CLONE=IMAGE:1927481 /UG=Hs.12421 hypothetical protein	AI348009	1.68
lactate dehydrogenase C	NM_002301	1.68
APG5 autophagy 5-like (S. cerevisiae)	AF293841	1.68
ectonucleoside triphosphate diphosphohydrolase 6 (putative function)	NM_001247	1.68
hypothetical protein MGC4022	AI805266	1.68
zinc finger protein 161 homolog (mouse)	NM_003409	1.68
ribosomal protein L12	AA281332	1.68
Consensus includes gb:AF064606.1 /DEF=Homo sapiens KB07 protein mRNA, partial cds. /FEA=mRNA /PROD=KB07 protein /DB_XREF=gi:3152665 /UG=Hs.306242 Homo sapiens KB07 pr	AF064606	1.68
excision repair cross-complementing rodent repair deficiency, complementation group 1 (includes overlapping antisense sequence)	NM_001983	1.68
lactate dehydrogenase A	NM_005566	1.68
hypothetical protein FLJ22490	BE044503	1.68
golgi associated, gamma adaptin ear containing, ARF binding protein 2	AA868560	1.68
leucine-rich PPR-motif containing	AI653608	1.68
fatty-acid-Coenzyme A ligase, long-chain 2	NM_021122	1.68
hypothetical protein MGC2835	NM_024072	1.68
interferon, alpha 21	M12350	1.67
prostaglandin D2 synthase, hematopoietic	NM_014485	1.67
GNAS complex locus	AA401492	1.67
eIEF associated protein HSPC021	NM_016091	1.67
membrane cofactor protein (CD46, trophoblast-lymphocyte cross-reactive antigen)	D84105	1.67
chaperonin containing TCP1, subunit 6A (zeta 1)	NM_001762	1.67
glia maturation factor, beta	BC005359	1.67
BCL2-associated X protein	U19599	1.67
putative mitochondrial outer membrane protein import receptor	AB019219	1.67

hypothetical protein FLJ13902	NM_024653	1.67
CCAAT/enhancer binding protein (C/EBP), beta	AL564683	1.67
protein kinase, DNA-activated, catalytic polypeptide	U47077	1.67
steroid dehydrogenase homolog	NM_016142	1.67
8D6 antigen	NM_016579	1.67
Consensus includes gb:AW371924 /FEA=EST /DB_XREF=gi:6876578 /DB_XREF=est:RC4-BT0311-251199-012-a07 /UG=Hs.313848 ESTs, Moderately similar to MGN_HUMAN MAGO NASHI	AW371924	1.67
hypothetical protein FLJ13149	NM_021826	1.67
osteoclast stimulating factor 1	NM_012383	1.67
three prime repair exonuclease 1	NM_016381	1.67
SRB7 suppressor of RNA polymerase B homolog (yeast)	AI688580	1.67
nucleoporin 50kD	NM_007172	1.67
frizzled homolog 3 (Drosophila)	NM_017412	1.67
nucleoporin 155kD	NM_004298	1.67
chromosome 21 open reading frame 66	NM_016631	1.67
hemoglobin, zeta	NM_005332	1.67
inositol 1,4,5-triphosphate receptor, type 3	NM_002224	1.67
delta-like 3 (Drosophila)	NM_016941	1.67
mitogen-activated protein kinase 1	AA195999	1.67
hypothetical protein MGC5618	BF575213	1.67
X-ray repair complementing defective repair in Chinese hamster cells 5 (double-strand-break rejoining; Ku autoantigen, 80kD)	AA205834	1.67
RAS protein activator like 1 (GAP1 like)	NM_004658	1.67
FK506 binding protein 12-rapamycin associated protein 1	U88966	1.67
LPAP for lysophosphatidic acid phosphatase	NM_016361	1.67
RAB35, member RAS oncogene family	BF791960	1.66
cathepsin C	NM_001814	1.66
glucocorticoid modulatory element binding protein 1	NM_006582	1.66
numb homolog (Drosophila)	AF015040	1.66
hypothetical protein FLJ22222	AI885411	1.66
TERF1 (TRF1)-interacting nuclear factor 2	NM_012461	1.66
tumor protein p53-binding protein	AI150117	1.66
gb:BC001121.1 /DEF=Homo sapiens, Similar to deoxyguanosine kinase, clone MGC:2111, mRNA, complete cds. /FEA=mRNA /PROD=Similar to deoxyguanosine kinase /DB_XREF=gi:12654	BC001121	1.66
pituitary tumor-transforming 1	NM_004219	1.66
chaperonin containing TCP1, subunit 6B (zeta 2)	NM_006584	1.66
putative tumor suppressor	AF040704	1.66
tumor necrosis factor receptor superfamily, member 6	X83493	1.66
mitofusin 1	AF329637	1.66
PDGFA associated protein 1	NM_014891	1.66
hypothetical protein FLJ22584	NM_024525	1.66
hypothetical protein FLJ12118	NM_024537	1.66
myosin, heavy polypeptide 9, non-muscle	AI827941	1.66
BCL2-like 1	NM_001191	1.66
hypothetical protein bK1048E9.5	AL041451	1.66
single-stranded DNA-binding protein 1	NM_003143	1.66
hypothetical protein FLJ21610	NM_022751	1.66
Cluster Incl. AA150503:zl08g04.s1 Homo sapiens cDNA, 3' end /clone=IMAGE-491766 /clone_end=3' /gb=AA150503 /gi=1722017 /ug=Hs.24435 /len=631	AA150503	1.66
solute carrier family 16 (monocarboxylic acid transporters), member 3	AL513917	1.66
Link guanine nucleotide exchange factor II	NM_016339	1.66
isocitrate dehydrogenase 3 (NAD+) gamma	U69268	1.66
U2(RNU2) small nuclear RNA auxillary factor 1 (non-standard symbol)	NM_006758	1.66
dihydrolipoamide dehydrogenase (E3 component of pyruvate dehydrogenase complex, 2-oxo-glutarate complex, branched chain keto acid dehydrogenase complex)	J03620	1.66
Consensus includes gb:AV700332 /FEA=EST /DB_XREF=gi:10302303 /DB_XREF=est:AV700332 /CLONE=GKCDIF11 /UG=Hs.243901 Homo sapiens cDNA FLJ20738 fis, clone HEP08257	AV700332	1.66
CGI-96 protein	Z93241	1.66
RAS guanyl releasing protein 2 (calcium and DAG-regulated)	AI688812	1.66
toll-like receptor 7	NM_016562	1.66
hypothetical protein FLJ20558	NM_017880	1.66
hypothetical protein FLJ11896	NM_024611	1.66
zinc finger protein 212	NM_012256	1.66
nuclear respiratory factor 1	NM_005011	1.66
cyclin-dependent kinase (CDC2-like) 10	NM_003674	1.66
EH-domain containing 4	AF320070	1.66
death-associated protein 6	NM_001350	1.66
uncharacterized bone marrow protein BM029	NM_018450	1.66
hypothetical protein FLJ20530	NM_017864	1.66

cholinergic receptor, nicotinic, alpha polypeptide 1 (muscle)	NM_000079	1.65
ATP synthase, H+ transporting, mitochondrial F0 complex, subunit g	AF070655	1.65
BRF1 homolog, subunit of RNA polymerase III transcription initiation factor IIIB (S.cerevisiae)	N91109	1.65
SON DNA binding protein	X63071	1.65
glutamyl-prolyl-tRNA synthetase	AI475965	1.65
signal transducer and activator of transcription 3 (acute-phase response factor)	AA634272	1.65
spleen tyrosine kinase	NM_003177	1.65
GDP-mannose pyrophosphorylase B	NM_021971	1.65
purinergic receptor P2X, ligand-gated ion channel, 7	NM_002562	1.65
DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 18 (Myc-regulated)	BC003360	1.65
protein kinase, AMP-activated, gamma 1 non-catalytic subunit	NM_002733	1.65
protein tyrosine kinase 9-like (A6-related protein)	NM_007284	1.65
capping protein (actin filament) muscle Z-line, beta	AL035413	1.65
LIN-7 protein 3	NM_018362	1.65
deleted in lung and esophageal cancer 1	AI860341	1.65
ubiquinol-cytochrome c reductase complex (7.2 kD)	NM_013387	1.65
mutS homolog 6 (E. coli)	D89646	1.65
hypothetical protein FLJ10143	NM_018009	1.65
ubiquinol-cytochrome c reductase (6.4kD) subunit	NM_006830	1.65
DKFZP586I0619 protein	BF055496	1.65
hypothetical protein	NM_019083	1.65
Sec61 gamma	NM_014302	1.65
heterogeneous nuclear ribonucleoprotein K	BC000355	1.65
replication protein A2 (32kD)	NM_002946	1.65
Consensus includes gb:BE733979 /FEA=EST /DB_XREF=gi:10147971 /DB_XREF=est:601571007F1 /CLONE=IMAGE:3925800 /UG=Hs.82202 ribosomal protein L17	BE733979	1.65
BRCA2 and CDKN1A interacting protein	NM_016567	1.65
fON3 protein	AB019494	1.65
LIM protein (similar to rat protein kinase C-binding enigma)	NM_006457	1.65
ribosomal protein L13	BC004954	1.65
zinc finger protein 42 (myeloid-specific retinoic acid- responsive)	NM_003422	1.65
HCGVIII-1 protein	X92110	1.64
amyloid beta (A4) precursor-like protein 2	AI525212	1.64
KIAA0332 protein	AU152088	1.64
calumenin	U67280	1.64
vacuolar protein sorting 45B (yeast)	AF165513	1.64
KIAA0699 protein	N45111	1.64
KIAA1491 protein	AK026088	1.64
hepatoma-derived growth factor (high-mobility group protein 1-like)	NM_004494	1.64
c-Mpl binding protein	AI743740	1.64
general transcription factor IIF, polypeptide 1 (74kD subunit)	NM_002096	1.64
Consensus includes gb:T33068 /FEA=EST /DB_XREF=gi:615166 /DB_XREF=est:EST56576 /UG=Hs.7101 anaphase-promoting complex subunit 5	T33068	1.64
solute carrier family 35 (CMP-sialic acid transporter), member 1	NM_006416	1.64
ribosomal protein S4, X-linked	AW132023	1.64
excision repair cross-complementing rodent repair deficiency, complementation group 3 (xeroderma pigmentosum group B complementing)	NM_000122	1.64
GDP-mannose 4,6-dehydratase	AI762113	1.64
KIAA0350 protein	AA731693	1.64
KIAA0365 gene product	AI928127	1.64
GTP cyclohydrolase 1 (dopa-responsive dystonia)	NM_000161	1.64
tumor protein D52	BE974098	1.64
hypothetical protein FLJ21877	AI286226	1.64
DKFZP564K2062 protein	NM_015421	1.64
tripartite motif-containing 28	NM_005762	1.64
squalene epoxidase	AF098865	1.64
KIAA0020 gene product	NM_014878	1.64
chromosome 11 hypothetical protein ORF3	NM_020154	1.64
KIAA0476 gene product	NM_014856	1.64
RAN, member RAS oncogene family	BF112006	1.64
tubulin, gamma 1	NM_001070	1.64
hypothetical protein FLJ13194	NM_025146	1.64
methyl-Cp6 binding domain protein 2	AF072242	1.64
peroxisome biogenesis factor 10	BF339133	1.64
hypothetical protein	NM_016458	1.64
H2B histone family, member G	NM_003522	1.64

CGI-203 protein	NM_020408	1.64
serine/threonine kinase 25 (STE20 homolog, yeast)	NM_006374	1.64
esterase D/formylglutathione hydrolase	BC001169	1.63
chondroitin sulfate proteoglycan BEHAB/brevican	NM_021948	1.63
hypothetical protein FLJ22116	NM_024624	1.63
hypothetical protein FLJ20550	NM_017874	1.63
myotubularin related protein 4	AB014547	1.63
mitochondrial ribosomal protein L16	NM_017840	1.63
SnRNP assembly defective 1 homolog	NM_006590	1.63
actin related protein 2/3 complex, subunit 1B (41 kD)	NM_005720	1.63
RAS p21 protein activator (GTPase activating protein) 1	M23612	1.63
hypothetical protein FLJ21016	NM_025160	1.63
cytochrome c oxidase subunit VIIc pseudogene 1	AF042165	1.63
receptor (TNFRSF)-interacting serine-threonine kinase 1	U50062	1.63
AFG3 ATPase family gene 3-like 2 (yeast)	NM_006796	1.63
presenilins associated rhomboid-like protein	NM_018622	1.63
HSPC028 protein	NM_014038	1.63
Consensus includes gb:BF793446 /FEA=EST /DB_XREF=gi:12098500 /DB_XREF=est:602254922F1 /CLONE=IMAGE:4347388 /UG=Hs.123122 FSH primary response (LRPR1, rat) homolog	BF793446	1.63
alcohol dehydrogenase 5 (class III), chi polypeptide	M29872	1.63
similar to Caenorhabditis elegans protein C42C1.9	NM_006459	1.63
glutamate receptor, ionotrophic, AMPA 3	NM_000828	1.63
Consensus includes gb:AU154486 /FEA=EST /DB_XREF=gi:11016007 /DB_XREF=est:AU154486 /CLONE=NT2RP4001238 /UG=Hs.119023 SMC2 (structural maintenance of chromosomes ;	AU154486	1.63
KIAA1012 protein	NM_014939	1.63
protein tyrosine phosphatase, receptor type, C	NM_002838	1.63
syntaxin 7	NM_003569	1.63
HUS1 checkpoint homolog (S. pombe)	NM_004507	1.63
AND-1 protein	AW772140	1.63
TGFB1-induced anti-apoptotic factor 1	NM_004740	1.63
KIAA0194 protein	D83778	1.63
metallothionein IV	X55503	1.63
solute carrier family 29 (nucleoside transporters), member 1	AF079117	1.63
SH2 domain protein 1A, Duncan's disease (lymphoproliferative syndrome)	AF072930	1.63
HEMK homolog 7kb	AF131220	1.63
putative glioblastoma cell differentiation-related	NM_016172	1.63
hypothetical protein MGC2494	NM_023933	1.63
apoptosis inhibitor 5	AF229253	1.63
tropomyosin 4	AI214061	1.63
thyroid hormone receptor interactor 12	NM_004238	1.63
ATP-binding cassette, sub-family G (WHITE), member 1	NM_004915	1.62
UBX domain-containing 2	AU149908	1.62
heterogeneous nuclear ribonucleoprotein A3	BF195526	1.62
hypothetical protein FLJ20897	BF690020	1.62
hypothetical protein FLJ12496	NM_022902	1.62
Consensus includes gb:BF115739 /FEA=EST /DB_XREF=gi:10985215 /DB_XREF=est:7n64b08.x1 /CLONE=IMAGE:3569246 /UG=Hs.15740 Homo sapiens mRNA; cDNA DKFZp434E033 (fro	BF115739	1.62
interferon, alpha 13	NM_006900	1.62
allantoicase	NM_018436	1.62
ubiquitin-conjugating enzyme E2L 3	BE964689	1.62
T-cell leukemia/lymphoma 6	NM_014418	1.62
hypothetical protein MGC2488	BC000229	1.62
similar to S. cerevisiae RER1	AF157324	1.62
myosin, light polypeptide 4, alkali; atrial, embryonic	AF116676	1.62
cytosolic acyl coenzyme A thioester hydrolase	AL031848	1.62
ATP-binding cassette, sub-family B (MDR/TAP), member 8	NM_007188	1.62
survival of motor neuron protein interacting protein 1	AB037702	1.62
non-POU-domain-containing, octamer-binding	NM_007363	1.62
hypothetical protein MGC3067	NM_024295	1.62
paired-like homeodomain transcription factor 1	U70370	1.62
gamma-glutamyl carboxylase	BE326952	1.62
B-cell CLL/lymphoma 7B	NM_001707	1.62
hypothetical protein FLJ12788	NM_022492	1.62
nth endonuclease III-like 1 (E. coli)	AC005600	1.62
putative membrane protein	AF274935	1.62
KIAA0801 gene product	NM_014829	1.62

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homolog of Yeast RRP4 (ribosomal RNA processing 4), 3'-5'-exoribonuclease	NM_014285	1.62
hepatocyte growth factor-regulated tyrosine kinase substrate	AF260566	1.62
frequently rearranged in advanced T-cell lymphomas 2	AB045118	1.62
protein tyrosine phosphatase, non-receptor type 6	NM_002831	1.62
Consensus includes gb:BE568184 /FEA=EST /DB_XREF=gi:9811904 /DB_XREF=est:601341722F1 /CLONE=IMAGE:3683860 /UG=Hs.200400 Human DNA sequence from BAC 15E1 on chro	BE568184	1.62
hypothetical protein PRO1843	BF247371	1.62
tubulin, alpha, ubiquitous	BC006379	1.62
TATA-binding protein-binding protein	NM_013375	1.62
stearoyl-CoA desaturase (delta-9-desaturase)	AB032261	1.62
SRp25 nuclear protein	NM_016638	1.62
oxysterol binding protein	AI927993	1.62
general transcription factor IIE, polypeptide 2 (beta subunit, 34kD)	NM_002095	1.61
hypothetical protein FLJ12547	NM_024992	1.61
KIAA0912 protein	AB020719	1.61
hypothetical protein FLJ20550	AI761506	1.61
polymerase (DNA-directed) sigma	NM_006999	1.61
hypothetical protein	NM_019049	1.61
pre-B lymphocyte gene 3	NM_013378	1.61
putative membrane protein	AB020980	1.61
fms-related tyrosine kinase 3 ligand	NM_001459	1.61
transmembrane protein 1	NM_003274	1.61
a disintegrin and metalloproteinase domain 17 (tumor necrosis factor, alpha, converting enzyme)	AI797833	1.61
Cp6 binding protein	U55777	1.61
dynamin 1-like	NM_012062	1.61
gb:NM_016473.1 /DEF=Homo sapiens hypothetical protein (LOC51243), mRNA. /FEA=CDS /GEN=LOC51243 /PROD=hypothetical protein /DB_XREF=gi:7706014 /UG=Hs.267263 hypothet	NM_016473	1.61
Consensus includes gb:BE780075 /FEA=EST /DB_XREF=gi:10201273 /DB_XREF=est:601469954F1 /CLONE=IMAGE:3873157 /UG=Hs.74137 transmembrane trafficking protein	BE780075	1.61
hippocalcin-like 1	BE617588	1.61
insulin induced protein 2	AL080184	1.61
glucocorticoid receptor DNA binding factor 1	AI670100	1.61
glucose-6-phosphatase, catalytic (glycogen storage disease type I, von Gierke disease)	AF097831	1.61
eukaryotic translation initiation factor 2, subunit 2 (beta, 38kD)	AL031668	1.61
cyclin T2	NM_001241	1.61
ubiquitin-conjugating enzyme E2 variant 1	BG164064	1.61
protein similar to E.coli yhdg and R. capsulatus nifR3	AW242981	1.61
nucleolin	NM_005381	1.61
v-myb myeloblastosis viral oncogene homolog (avian)-like 1	AW592266	1.60
hypothetical protein FLJ20986	NM_024524	1.60
hydroxyacyl-Coenzyme A dehydrogenase/3-ketoacyl-Coenzyme A thiolase/enoyl-Coenzyme A hydratase (trifunctional protein), alpha subunit	BG472176	1.60
KIAA0728 protein	AI798790	1.60
gb:BC002361.1 /DEF=Homo sapiens, Similar to transducin (beta)-like 3, clone MGC:8613, mRNA, complete cds. /FEA=mRNA /PROD=Similar to transducin (beta)-like 3 /DB_XREF=gi:1280	BC002361	1.60
platelet-activating factor acetylhydrolase, isoform Ib, alpha subunit (45kD)	L13387	1.60
KIAA0978 protein	N64780	1.60
serine/threonine kinase 19	NM_004197	1.60
staufen, RNA binding protein (Drosophila)	AI132258	1.60
Consensus includes gb:AV701318 /FEA=EST /DB_XREF=gi:10717648 /DB_XREF=est:AV701318 /CLONE=ADAAGD10 /UG=Hs.278626 ArgAbl-interacting protein ArgBP2	AV701318	1.60
prolactin regulatory element binding	NM_013388	1.60
bridging integrator 3	AK001289	1.60
transcription elongation factor B (SIII), polypeptide 1 (15kD, elongin C)	NM_005648	1.60
polymerase (DNA directed), epsilon 2	NM_002692	1.60
CCR4-NOT transcription complex, subunit 2	BG105204	1.60
STIP1 homology and U-Box containing protein 1	NM_005861	1.60
lactate dehydrogenase B	NM_002300	1.60
hypothetical protein FLJ14888	AF161382	1.60
hypothetical protein MGC5466	AI589507	1.60
splicing factor, arginine/serine-rich 6	AL031681	1.60
chloride channel, nucleotide-sensitive, 1A	AF005422	1.60
dUTP pyrophosphatase	AB049113	1.60
protein tyrosine kinase 9	AW665024	1.60
glucose-6-phosphate dehydrogenase	NM_000402	1.60
chaperonin containing TCP1, subunit 2 (beta)	NM_006431	1.60
solute carrier family 35 (UDP-N-acetylglucosamine (UDP-GlcNAc) transporter), member 3	NM_012243	1.60
nuclear localization signal deleted in velocardiofacial syndrome	NM_003776	1.60

beclin 1 (coiled-coil, myosin-like BCL2 interacting protein)	AF139131	1.60
polycystic kidney disease 1 (autosomal dominant)	NM_000296	1.60
RAP1B, member of RAS oncogene family	NM_015646	1.59
hypothetical protein FLJ12525	NM_031206	1.59
hypothetical protein FLJ10482	NM_018107	1.59
hypothetical protein FLJ10486	NM_018109	1.59
putative c-Myc-responsive	AF040105	1.59
protein kinase D2	AL050147	1.59
ribosomal protein S20	NM_001023	1.59
GLE1 RNA export mediator-like (yeast)	NM_001499	1.59
Rab9 effector p40	NM_005833	1.59
NICE-5 protein	NM_017582	1.59
neuroblastoma-amplified protein	NM_015909	1.59
hypothetical protein FLJ20640	BC000890	1.59
inositol polyphosphate-5-phosphatase, 40kD	NM_005539	1.59
FtsJ homolog 1 (E. coli)	AV723177	1.59
hypothetical protein FZ5965	BF215644	1.59
karyopherin alpha 3 (importin alpha 4)	AF034756	1.59
MLN51 protein	NM_007359	1.59
epoxide hydrolase 2, cytoplasmic	AF233336	1.59
G-rich RNA sequence binding factor 1	NM_002092	1.59
KIAA0943 protein	AA532726	1.59
vascular endothelial growth factor B	NM_003377	1.59
EH-domain containing 1	AW182860	1.59
TED protein	NM_015686	1.59
centromere protein A (17kD)	BC002703	1.59
hydroxyacyl-Coenzyme A dehydrogenase/3-ketoacyl-Coenzyme A thiolase/enoyl-Coenzyme A hydratase (trifunctional protein), beta subunit	NM_000183	1.59
mitochondrial ribosomal protein S27	D87453	1.59
histidine triad nucleotide binding protein	U27143	1.59
KIAA0286 protein	AW205215	1.59
solute carrier family 2 (facilitated glucose/fructose transporter), member 5 -.	NM_003039	1.59
mitochondrial translational release factor 1	NM_004294	1.59
X-ray repair complementing defective repair in Chinese hamster cells 4	AB017445	1.59
ubiquitination factor E4B (UFD2 homolog, yeast)	AF091093	1.59
SCO cytochrome oxidase deficient homolog 2 (yeast)	NM_005138	1.59
platelet-activating factor acetylhydrolase, isoform Ib, alpha subunit (45kD)	BE256969	1.59
ITV1 gene	AI928526	1.59
coxsackie virus and adenovirus receptor	NM_001338	1.59
DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 19 (DBP5 homolog, yeast)	AL553254	1.59
surfeit 5	NM_006752	1.59
Cluster Incl. AL050297:Homo sapiens mRNA; cDNA DKFZp564N123 (from clone DKFZp564N123) /cds=UNKNOWN /gb=AL050297 /gi=4886448 /ug=Hs.25894 /len=1340	AL050297	1.59
mitochondrial ribosomal protein L15	NM_014175	1.59
BAF53	NM_004301	1.59
GS2 gene	U03886	1.59
signal transducing adaptor molecule (SH3 domain and ITAM motif) 1	NM_003473	1.59
mitochondrial ribosomal protein L35	NM_016622	1.59
ariadne homolog, ubiquitin-conjugating enzyme E2 binding protein, 1 (Drosophila)	NM_005744	1.59
B-cell CLL/lymphoma 6 (zinc finger protein 51)	NM_001706	1.59
hypothetical protein	NM_019023	1.59
actin, gamma 1	BE741683	1.59
CGI-26 protein	NM_015954	1.59
putative T1/ST2 receptor binding protein	NM_006858	1.59
v-rel reticuloendotheliosis viral oncogene homolog B, nuclear factor of kappa light polypeptide gene enhancer in B-cells 3 (avian)	NM_006509	1.59
ClpP caseinolytic protease, ATP-dependent, proteolytic subunit homolog (E. coli)	NM_006012	1.59
protease, serine, 22	NM_022119	1.59
coilin	BG391060	1.59
protein kinase domains containing protein similar to phosphoprotein C8FW	NM_021158	1.59
Consensus includes gb:AW450910 /FEA=EST /DB_XREF=gi:6991686 /DB_XREF=est:UI-H-BI3-all-e-01-0-UI.s1 /CLONE=IMAGE:2737225 /UG=Hs.74101 spleen tyrosine kinase /FL=gb:BC0	AW450910	1.59
A kinase (PRKA) anchor protein 13	M90360	1.59
CGI-100 protein	NM_016040	1.59
ribosomal protein L30	L05095	1.58
interleukin 16 (lymphocyte chemoattractant factor)	NM_004513	1.58
KIAA0409 protein	NM_015324	1.58

hypothetical protein FLJ22405	AY007098	1.58
SRB7 suppressor of RNA polymerase B homolog (yeast)	U46837	1.58
KIAA0842 protein	AB020649	1.58
cargo selection protein (mannose 6 phosphate receptor binding protein)	NM_005817	1.58
nuclear factor of activated T-cells, cytoplasmic, calcineurin-dependent 1	U80918	1.58
microspherule protein 1	NM_006337	1.58
hypothetical protein FLJ10290	NM_018047	1.58
elastase 3A, pancreatic (protease E)	BC005918	1.58
cysteinyl-tRNA synthetase	NM_001751	1.58
putative methyltransferase	BC001650	1.58
Consensus includes gb:AL133228 /DEF=Human DNA sequence from clone RP5-1071L10 on chromosome 20 Contains part of a gene for a new member of the thymosininterferon-inducible m	AL133228	1.58
hypothetical protein FLJ20568	AA436930	1.58
hypothetical protein FLJ10241	AI768100	1.58
thimet oligopeptidase 1	NM_003249	1.58
actin related protein 2/3 complex, subunit 2 (34 kD)	AF116702	1.58
Gp1-interacting zinc finger protein	NM_012127	1.58
steroid sulfatase (microsomal), arylsulfatase C, isozyme S	NM_000351	1.58
hypothetical protein FLJ12553	NM_024700	1.58
zinc metalloproteinase (STE24 homolog, yeast)	NM_005857	1.58
thiosulfate sulfurtransferase (rhodanese)	D87292	1.58
protein kinase, interferon-inducible double stranded RNA dependent activator	AF083033	1.58
partner of RAC1 (arfaptin 2)	NM_012402	1.58
ubiquitin specific protease 14 (tRNA-guanine transglycosylase)	NM_005151	1.58
activating transcription factor 4 (tax-responsive enhancer element B67)	NM_001675	1.58
stromal antigen 2	NM_006603	1.58
KIAA1856 protein	AL080102	1.58
hypothetical protein FLJ20011	BC003086	1.58
p47 protein	NM_018839	1.58
DKFZP564M082 protein	NM_014042	1.58
nuclear receptor subfamily 1, group D, member 1	X72631	1.58
transducin-like enhancer of split 4 (E(sp1) homolog, Drosophila)	AL358975	1.58
solute carrier family 25 (mitochondrial carrier; oxoglutarate carrier), member 11	NM_003562	1.58
solute carrier family 16 (monocarboxylic acid transporters), member 1	AL162079	1.58
hypothetical protein FLJ10407	NM_018087	1.58
progesterone receptor membrane component 2	NM_006320	1.58
second mitochondria-derived activator of caspase	NM_019887	1.58
ribosomal protein S17	BF026595	1.58
H4 histone family, member E	NM_021968	1.58
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ferritin, heavy polypeptide pseudogene 1	J04755	1.58
Ras homolog enriched in brain 2	NM_005614	1.58
hypothetical protein FLJ12595	NM_024994	1.58
Consensus includes gb:BG105365 /FEA=EST /DB_XREF=gi:12599141 /DB_XREF=est:602312473F1 /CLONE=IMAGE:4422043 /UG=Hs.23348 S-phase kinase-associated protein 2 (p45) /F	BG105365	1.58
translational inhibitor protein p14.5	N54448	1.57
polo (Drosophila)-like kinase	NM_005030	1.57
protein phosphatase 2A, regulatory subunit B' (PR 53)	BC002545	1.57
MAGUK protein p55T; Protein Associated with Lins 2	NM_016447	1.57
NADH dehydrogenase (ubiquinone) Fe-S protein 5 (15kD) (NADH-coenzyme Q reductase)	NM_004552	1.57
Consensus includes gb:AA203487 /FEA=EST /DB_XREF=gi:1799460 /DB_XREF=est:zx53d03.r1 /CLONE=IMAGE:446213 /UG=Hs.314363 ESTs	AA203487	1.57
calumenin	BF939365	1.57
DKFZP566C0424 protein	BG171020	1.57
UDP-Gal:betaGlcNAc beta 1,4- galactosyltransferase, polypeptide 4	BC004523	1.57
KIAA0009 gene product	BF214329	1.57
guanine monphosphate synthetase	NM_003875	1.57
inositol polyphosphate-4-phosphatase, type I, 107kD	NM_004027	1.57
integrin, beta 1 (fibronectin receptor, beta polypeptide, antigen CD29 includes MDF2, MSK12)	AA215854	1.57
phosphoglucomutase 1	NM_002633	1.57
survival of motor neuron protein interacting protein 1	AB037703	1.57
peroxisomal biogenesis factor 6	D83703	1.57
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polymyositis/scleroderma autoantigen 2 (100kD)	NM_002685	1.57
HIV-1 Tat interactive protein, 60 kDa	BC000166	1.57
SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily a, member 2	AV725365	1.57

beta-1,3-glucuronyltransferase 3 (glucuronosyltransferase I)	NM_012200	1.57
hypothetical protein FLJ22477	NM_024735	1.57
BCL2/adenovirus E1B 19kD interacting protein 3	NM_004052	1.57
hypothetical protein FLJ20719	AB051480	1.57
enoyl Coenzyme A hydratase 1, peroxisomal	NM_001398	1.57
DKFZP586J0119 protein	AJ011307	1.57
developmentally regulated GTP binding protein 1	NM_004147	1.57
mitochondrial ribosomal protein L46	NM_022163	1.57
pyrophosphatase (inorganic)	NM_021129	1.57
gb:NM_024030.1 /DEF=Homo sapiens hypothetical protein MGC3260 (MGC3260), mRNA. /FEA=mRNA /GEN=MGC3260 /PROD=hypothetical protein MGC3260 /DB_XREF=gi:13128977 /UG:	NM_024030	1.57
phosphofructokinase, liver	BC006422	1.57
calcineurin binding protein 1	AB002328	1.57
DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 5 (RNA helicase, 68kD)	NM_004396	1.57
STE20-like kinase	NM_016281	1.57
KIAA0370 protein	AL546600	1.57
KDEL (Lys-Asp-Glu-Leu) endoplasmic reticulum protein retention receptor 2	AL542253	1.57
mannose-P-dolichol utilization defect 1	AF059752	1.57
RAD23 homolog A (S. cerevisiae)	NM_005053	1.57
hypothetical protein FLJ21212	NM_024642	1.57
ADP-ribosylation factor-like 6 interacting protein	D31885	1.56
protein phosphatase 1, regulatory (inhibitor) subunit 2	NM_006241	1.56
programmed cell death 5	NM_004708	1.56
elastase 3A, pancreatic (protease E)	D00306	1.56
hypothetical protein FLJ20561	NM_017882	1.56
ubiquitin-conjugating enzyme E2E 3 (UBC4/5 homolog, yeast)	AB017644	1.56
huntingtin interacting protein-1-related	AB014555	1.56
DnaJ (Hsp40) homolog, subfamily C, member 4	NM_005528	1.56
hypothetical protein FLJ10512	AL133215	1.56
DKFZP547E2110 protein	AB028628	1.56
centromere protein A (17kD)	NM_001809	1.56
phospholipase C, gamma 1 (formerly subtype 148)	AL022394	1.56
centaurin, beta 1	NM_014716	1.56
karyopherin (importin) beta 1	BC003572	1.56
kinesin-associated protein 3	NM_014970	1.56
Consensus includes gb:AL515874 /FEA=EST /DB_XREF=gi:12779367 /DB_XREF=est:AL515874 /CLONE=CS0DA001YF06 (3 prime) /UG=Hs.83086 Homo sapiens GT212 mRNA	AL515874	1.56
iduronate 2-sulfatase (Hunter syndrome)	NM_000202	1.56
WD repeat domain 13	AF158978	1.56
transmembrane 4 superfamily member 7	AF054841	1.56
lung cancer candidate	BF308548	1.56
2,4-dienoyl CoA reductase 1, mitochondrial	NM_001359	1.56
gb:U95006.1 /DEF=Human D9 splice variant A mRNA, complete cds. /FEA=mRNA /PROD=D9 splice variant A /DB_XREF=gi:2071992 /UG=Hs.37616 Human D9 splice variant B mRNA, compl	U95006	1.56
LanC (bacterial lantibiotic synthetase component C)-like 2	NM_018697	1.56
nucleoporin-like protein 1	NM_007342	1.56
hypothetical protein FLJ20477	AA203365	1.56
actinin, alpha 4	U48734	1.56
CDC5 cell division cycle 5-like (S. pombe)	AW268817	1.56
APG12 autophagy 12-like (S. cerevisiae)	BE965998	1.56
NS1-binding protein	AB020657	1.56
KIAA0246 protein	NM_015136	1.56
similar to vaccinia virus HindIII K4L ORF	NM_012268	1.56
Consensus includes gb:BE880828 /FEA=EST /DB_XREF=gi:10329604 /DB_XREF=est:601493261F1 /CLONE=IMAGE:3895407 /UG=Hs.84775 Human transposon-like element mRNA	BE880828	1.56
katanin p80 (WD40-containing) subunit B 1	NM_005886	1.56
hypothetical protein FLJ11749	NM_024591	1.56
acidic protein rich in leucines	NM_006401	1.56
regulator of G-protein signalling 14	AF037195	1.56
heterogeneous nuclear ribonucleoprotein U (scaffold attachment factor A)	BC003621	1.55
eukaryotic translation initiation factor 2B, subunit 3 (gamma, 58kD)	NM_020365	1.55
basigin (OK blood group)	AL550657	1.55
hypothetical protein FLJ20729	NM_017953	1.55
hypothetical protein FLJ13134	NM_024701	1.55
protein kinase C, nu	NM_005813	1.55
sorting nexin 3	AB047360	1.55
RAP2A, member of RAS oncogene family	AI302106	1.55

Nef-associated factor 1	NM_006058	1.55
loricrin	NM_000427	1.55
HIV-1 Tat interactive protein 2, 30 kDa	BC002439	1.55
hypothetical protein	NM_014886	1.55
Consensus includes gb:AL031669 /DEF=Human DNA sequence from clone RP4-661I20 on chromosome 20q11.23-12 Contains a gene for a protein similar to RPL12 (ribosomal protein L12),	AL031669	1.55
Cbp/p300-interacting transactivator, with Glu/Asp-rich carboxy-terminal domain, 2	NM_006079	1.55
peroxisomal biogenesis factor 14	AB017546	1.55
golgi SNAP receptor complex member 2	AW149492	1.55
ornithine decarboxylase antizyme 2	AF242521	1.55
methyl-CpG binding domain protein 3	NM_003926	1.55
S88I26 protein	NM_018846	1.55
hypothetical protein FLJ20548	BC001719	1.55
hypothetical protein FLJ10101	NM_024718	1.55
coronin, actin binding protein, 1C	BC002342	1.55
hypothetical protein MGC3035	AL040896	1.55
gene amplified in squamous cell carcinoma 1	AB037901	1.55
RA-regulated nuclear matrix-associated protein	NM_016448	1.55
surfeit 1	NM_003172	1.55
hypothetical protein FLJ20406	NM_017806	1.55
heterogeneous nuclear ribonucleoprotein A2/B1	NM_002137	1.55
hypothetical protein FLJ21940	AL049305	1.55
peroxisomal farnesylated protein	NM_002857	1.55
proteasome (prosome, macropain) subunit, beta type, 4	NM_002796	1.55
protein tyrosine kinase 2 beta	NM_004103	1.55
hypothetical protein FLJ10808	NM_018227	1.55
hypothetical protein FLJ20373	NM_017792	1.55
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paired box gene 8	NM_013952	1.55
glutamate-cysteine ligase, catalytic subunit	BF676980	1.55
nuclease sensitive element binding protein 1	X96666	1.55
putative GTP-binding protein similar to RAY/RAB1C	NM_006860	1.55
serine hydroxymethyltransferase 2 (mitochondrial)	AW190316	1.55
ceroid-lipofuscinosis, neuronal 5	AV700514	1.55
hypothetical protein	U69190	1.55
protein kinase C, iota	L18964	1.55
insulin-degrading enzyme	AA918442	1.54
Consensus includes gb:AF123462 /DEF=Homo sapiens BAC526N18 neurexin III gene, partial cds /FEA=CDS /DB_XREF=gi:4240565 /UG=Hs.247837 Homo sapiens BAC526N18 neurexin III ;	AF123462	1.54
eukaryotic translation initiation factor 2, subunit 1 (alpha, 35kD)	NM_004094	1.54
coatomer protein complex, subunit alpha	AI621079	1.54
DC13 protein	NM_020188	1.54
phosphatase and tensin homolog (mutated in multiple advanced cancers 1), pseudogene 1	AF023139	1.54
KIAA0729 protein	AB018272	1.54
iduronate 2-sulfatase (Hunter syndrome)	NM_006123	1.54
microfibrillar-associated protein 2	NM_017459	1.54
topoisomerase (DNA) III beta	D87012	1.54
major histocompatibility complex, class I-like sequence	NM_001531	1.54
H2A histone family, member A	NM_021052	1.54
regucalcin (senescence marker protein-30)	D31815	1.54
ribosomal protein S3A	NM_001006	1.54
glycerylate reductase/hydroxypyruvate reductase	AK024386	1.54
phospholipase A2-activating protein	AF145020	1.54
isovaleryl Coenzyme A dehydrogenase	AK022777	1.54
hypothetical protein FLJ12448	NM_022895	1.54
MCM4 minichromosome maintenance deficient 4 (S. cerevisiae)	AI936566	1.54
peroxisomal short-chain alcohol dehydrogenase	NM_021004	1.54
hypothetical protein MGC2463	NM_024070	1.54
cell division cycle 2-like 1 (PITSLRE proteins)	NM_001787	1.54
ubiquitin-conjugating enzyme E2 variant 2	U62136	1.54
polymerase (RNA) II (DNA directed) polypeptide C (33kD)	BC000409	1.54
immunoglobulin heavy constant mu	S55735	1.54
karyopherin (importin) beta 1	AA861608	1.54
sorting nexin 3	AF062483	1.54
LGN protein	NM_013296	1.54

DKFZp586F1019 protein	AL050102	1.54
Consensus includes gb:BF593594 /FEA=EST /DB_XREF=gi:11685918 /DB_XREF=est:7q12a08.x1 /CLONE=IMAGE:3697839 /UG=Hs.274440 Homo sapiens mRNA; cDNA DKFZp434I139 (fre	BF593594	1.54
adaptor-related protein complex 2, beta 1 subunit	NM_001282	1.54
hypothetical protein MGC4276 similar to C68198	NM_030940	1.54
myosin, light polypeptide 4, alkali; atrial, embryonic	M36172	1.54
H3 histone, family 3B (H3.3B)	AW138159	1.54
baculoviral IAP repeat-containing 5 (survivin)	AA648913	1.54
sulfotransferase family, cytosolic, 1A, phenol-preferring, member 1	NM_001055	1.54
Consensus includes gb:AI949687 /FEA=EST /DB_XREF=gi:5741997 /DB_XREF=est:wq13g04.x1 /CLONE=IMAGE:2471190 /UG=Hs.173638 transcription factor 7-like 2 (T-cell specific, HMG	AI949687	1.54
mutY homolog (E. coli)	NM_012222	1.54
excision repair cross-complementing rodent repair deficiency, complementation group 1 (includes overlapping antisense sequence)	NM_001983	1.54
putative neurotransmitter receptor	NM_003967	1.53
dynactin 4 (p62)	NM_016221	1.53
Consensus includes gb:AI344141 /FEA=EST /DB_XREF=gi:4081347 /DB_XREF=est:tc02d09.x1 /CLONE=IMAGE:2062673 /UG=Hs.229850 ESTs, Highly similar to G01251 Rar protein H.sapi	AI344141	1.53
KIAA0090 protein	AI143233	1.53
ADP-ribosylation factor 3	M33384	1.53
hypothetical protein FLJ10335	NM_018062	1.53
KIAA0237 gene product	NM_014747	1.53
melanoma antigen, family D, 1	AF217963	1.53
a disintegrin and metalloproteinase domain 10	NM_001110	1.53
filamin B, beta (actin binding protein 278)	AV712733	1.53
DKFZP586L0724 protein	AU158148	1.53
zinc finger protein 45 (a Kruppel-associated box (KRAB) domain polypeptide)	NM_003425	1.53
growth factor receptor-bound protein 2	L29511	1.53
sorting nexin 7	NM_015976	1.53
hypothetical protein FLJ20073	NM_017654	1.53
stearoyl-CoA desaturase (delta-9-desaturase)	BC005807	1.53
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human homolog of Drosophila Scribble	AI469403	1.53
immunoglobulin heavy constant mu	BG340670	1.53
hexokinase 1	NM_000188	1.53
hypothetical protein DC42	NM_030921	1.53
dipeptidyltpeptidase 8	NM_017743	1.53
HCDI protein	AI347090	1.53
DKFZP434D1335 protein	AL117499	1.53
ribosomal protein L39-like 1	L05096	1.53
polynucleotide kinase 3'-phosphatase	NM_007254	1.53
upstream transcription factor 2, c-fos interacting	NM_003367	1.53
E74-like factor 2 (ets domain transcription factor)	AF256223	1.53
hypothetical protein FLJ12700	NM_024910	1.53
hypothetical protein FLJ23320	NM_024672	1.53
ribosomal protein L7	NM_000971	1.53
succinate-CoA ligase, GDP-forming, beta subunit	AF131748	1.53
ubiquitin-conjugating enzyme E2N (UBC13 homolog, yeast)	NM_003348	1.52
KIAA0637 gene product	NM_014838	1.52
ubiquitin fusion degradation 1-like	BC001049	1.52
UDP-Gal:betaGlcNAc beta 1,4- galactosyltransferase, polypeptide 5	AL035683	1.52
golgi associated, gamma adaptin ear containing, ARF binding protein 2	BF196572	1.52
integrin, alpha L (antigen CD11A (p180), lymphocyte function-associated antigen 1; alpha polypeptide)	AC002310	1.52
cleavage stimulation factor, 3' pre-RNA, subunit 1, 50kD	NM_001324	1.52
putative DNA binding protein	AL523380	1.52
IDN3 protein	NM_015384	1.52
cytoplasmic polyadenylation element binding protein	NM_030594	1.52
protective protein for beta-galactosidase (galactosialidosis)	NM_000308	1.52
actin, beta	D50604	1.52
RAB2, member RAS oncogene family	M28213	1.52
hypothetical protein MGC2454	NM_025264	1.52
huntingtin (Huntington disease)	L12392	1.52
dystroglycan 1 (dystrophin-associated glycoprotein 1)	NM_004393	1.52
SWAP-70 protein	AI139569	1.52
hypothetical protein MGC10963	NM_030573	1.52
protein tyrosine phosphatase type IVA, member 2	AF208850	1.52
cortistatin	AB000263	1.52

hypothetical protein FLJ13912	NM_022770	1.52
RING1 and YY1 binding protein	W84482	1.52
zinc finger protein 216	AW471220	1.52
mannosyl (alpha-1,6-)-glycoprotein beta-1,2-N-acetylglucosaminyltransferase	AW103265	1.52
SKI-interacting protein	NM_012245	1.52
protein (peptidyl-prolyl cis/trans isomerase) NIMA-interacting 1	NM_006221	1.52
WD40 protein Ciao1	NM_004804	1.52
hypothetical protein FLJ20608	NM_017900	1.52
tissue specific transplantation antigen P35B	NM_003313	1.52
hemochromatosis	AF115264	1.52
HIV-1 Tat interactive protein, 60 kDa	NM_006388	1.52
BCL2-antagonist of cell death	U66879	1.52
hypothetical protein PRO1580	NM_018502	1.52
H2B histone family, member F	NM_021062	1.52
ITV1 gene	NM_006303	1.52
signal recognition particle 72kD	BE856385	1.51
general transcription factor IIF, polypeptide 1 (74kD subunit)	AW190445	1.51
ribosomal protein S24	BC000523	1.51
tyrosine kinase, non-receptor, 1	NM_003985	1.51
KIAA0244 protein	BF430956	1.51
gb:NM_005703.2 /DEF=Homo sapiens upstream regulatory element binding protein 1 (UREB1), mRNA. /FEA=CDS /GEN=UREB1 /PROD=upstream regulatory element binding protein 1 /DB	NM_005703	1.51
t-complex 1	BC000665	1.51
GTT1 protein	NM_020151	1.51
ubiquitin-protein isopeptide ligase (E3)	NM_014671	1.51
karyopherin (importin) beta 3	BC000947	1.51
DKFZP586J1624 protein	NM_015537	1.51
tubulin, beta polypeptide	NM_001069	1.51
LDL receptor adaptor protein	AA169780	1.51
hypothetical protein FLJ14153	NM_022736	1.51
proteasome (prosome, macropain) subunit, beta type, 5	BC004146	1.51
interferon, gamma-inducible protein 30	NM_006332	1.51
hypothetical protein FLJ20003	BC005212	1.51
SNARE protein	NM_006555	1.51
phosphoinositide-3-kinase, regulatory subunit 4, p150	BF740111	1.51
zinc finger protein 216	AF062347	1.51
hypothetical protein MGC10753	NM_016628	1.51
TcD37 homolog	AF123539	1.51
polymerase (RNA) II (DNA directed) polypeptide C (33kD)	AJ224143	1.51
VAMP (vesicle-associated membrane protein)-associated protein A (33kD)	AF154847	1.51
hypothetical protein FLJ20596	AF336851	1.51
tetratricopeptide repeat domain 4	NM_004623	1.51
potassium voltage-gated channel, Shal-related subfamily, member 1	NM_004979	1.51
glutathione synthetase	L42531	1.51
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DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 9 (RNA helicase A, nuclear DNA helicase II; leukophysin)	NM_001357	1.51
karyopherin beta 2b, transportin	BG258639	1.51
glycogen synthase 1 (muscle)	NM_002103	1.51
KIAA0286 protein	AB006624	1.51
heat shock 10kD protein 1 (chaperonin 10)	NM_002157	1.51
ribosomal protein S28	BC000354	1.51
sentrin-specific protease	NM_021627	1.51
TAK1-binding protein 2	AF241230	1.51
eukaryotic translation elongation factor 1 epsilon 1	NM_004280	1.51
hypothetical protein MGC3180	NM_024041	1.51
v-erb-b2 erythroblastic leukemia viral oncogene homolog 2, neuro/glioblastoma derived oncogene homolog (avian)	X03363	1.51
mitogen-activated protein kinase kinase 7	AW007458	1.51
SET domain and mariner transposase fusion gene	NM_006515	1.51
hypothetical protein FLJ12615 similar to membrane protein, palmitoylated 3 (MAGUK p55 subfamily member 5)	NM_022474	1.51
hypothetical protein FLJ11164	NM_018346	1.51
H4 histone family, member D	NM_003541	1.50
ataxia telangiectasia mutated (includes complementation groups A, C and D)	U82828	1.50
tubulin, alpha, ubiquitous	BC006481	1.50
KIAA0999 protein	AB023216	1.50

CDC16 cell division cycle 16 homolog (S. cerevisiae)	AF164598	1.50
DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 16	N36997	1.50
hypoxia-inducible factor 1, alpha subunit (basic helix-loop-helix transcription factor)	NM_001530	1.50
proteasome (prosome, macropain) 26S subunit, non-ATPase, 1	NM_002807	1.50
hypothetical protein FLJ10583	NM_018148	1.50
hypothetical protein FLJ10814	NM_018230	1.50
KIAA0399 protein	NM_015113	1.50
hypothetical protein similar to beta-transducin family	NM_018096	1.50
dCMP deaminase	BC001286	1.50
metazin 1	BC001906	1.50
inhibitor of DNA binding 2, dominant negative helix-loop-helix protein	NM_002166	1.50
hypothetical protein MGC2745	NM_024117	1.50
Consensus includes gb:AW138902 /FEA=EST /DB_XREF=gi:6143220 /DB_XREF=est:UI-H-BI1-aeq-e-08-0-UI.s1 /CLONE=IMAGE:2720342 /UG=Hs.173108 Homo sapiens cDNA: FLJ21897 fi	AW138902	1.50
calcium/calmodulin-dependent protein kinase I	NM_003656	1.50
mitochondrial import receptor Tom22	NM_020243	1.50
KIAA0998 protein	AB023215	1.50
NAD kinase	AI334128	1.50
cadherin, EGF LAG seven-pass G-type receptor 1 (flamingo homolog, Drosophila)	BC000059	1.50
v-raf murine sarcoma 3611 viral oncogene homolog 1	NM_001654	1.50
RAN binding protein 2	D42063	1.50
tyrosine 3-monooxygenase/tryptophan 5-monooxygenase activation protein, beta polypeptide	BC001359	1.50
coated vesicle membrane protein	NM_006815	1.50
alanine-glyoxylate aminotransferase (oxalosis I; hyperoxaluria I; glycolicaciduria; serine-pyruvate aminotransferase)	D13368	1.50
RAB6B, member RAS oncogene family	BC002510	1.50
hypothetical protein MGC10940	BC004331	1.50
Down syndrome critical region gene 2	NM_003720	1.50
hypothetical protein	NM_015705	1.50
KIAA0515 protein	AK026025	1.50
NG23 protein	NM_025259	1.50
mitogen-activated protein kinase kinase kinase 7	NM_003188	1.50
ets variant gene 5 (ets-related molecule)	BF060791	1.50
Consensus includes gb:AI224869 /DEF=Homo sapiens CXCR4 gene encoding receptor CXCR4 /FEA=mRNA /DB_XREF=gi:3059119 /UG=Hs.89414 chemokine (C-X-C motif), receptor 4 (fusin	AI224869	1.50
chromobox homolog 1 (HP1 beta homolog Drosophila)	NM_006807	1.50
nucleoporin 62kD	NM_012346	1.50
programmed cell death 2	NM_002598	1.50
hypothetical protein FLJ12171	NM_024619	1.50
CD22 antigen	X59350	1.50
adenylate kinase 2	AL513611	1.50
karyopherin (importin) beta 1	BG545463	1.50
histidine triad nucleotide binding protein	N32864	1.50
acidic protein rich in leucines	AV712577	1.50
KIAA0769 gene product	NM_014824	1.50
E2F transcription factor 1	M96577	1.50
KIAA1886 protein	AA868332	1.50
eukaryotic translation initiation factor 4A, isoform 2	NM_001967	1.50
hypothetical protein FLJ22494	NM_024815	1.50
lysophospholipase I	AF077198	1.50
DEAD/H (Asp-Glu-Ala-Asp/His) box binding protein 1	NM_016166	1.50
hypothetical protein FLJ23309	NM_024896	1.50
myxovirus (influenza) resistance 1, homolog of murine (interferon-inducible protein p78)	NM_002462	1.50
KIAA1194 protein	NM_015455	1.49
glutaryl-Coenzyme A dehydrogenase	NM_013976	1.49
hypothetical protein FLJ13660 similar to CDK5 activator-binding protein C53	NM_025197	1.49
KIAA0056 protein	AI796581	1.49
16.7Kd protein	NM_016139	1.49
pancreatic polypeptide	M15788	1.49
gb:AF308302.1 /DEF=Homo sapiens serologically defined breast cancer antigen NY-BR-96 mRNA, complete cds. /FEA=mRNA /PROD=serologically defined breast cancer antigenNY-BR-96 /	AF308302	1.49
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nascent-polypeptide-associated complex alpha polypeptide	NM_005594	1.49
RAD50 homolog (S. cerevisiae)	NM_005732	1.49
biliverdin reductase B (flavin reductase (NADPH))	NM_000713	1.49
speckle-type POZ protein	NM_003563	1.49
laminin, alpha 3 (nicein (150kD), kalinin (165kD), BM600 (150kD), epilegrin)	X77598	1.49

RNA helicase family	AA156961	1.49
chromobox homolog 3 (HP1 gamma homolog, Drosophila)	BE748755	1.49
nuclear receptor co-repressor 2	NM_006312	1.49
golgin-67	AF163441	1.49
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signal recognition particle 72kD	AF069765	1.49
CDC16 cell division cycle 16 homolog (S. cerevisiae)	AF164598	1.49
ribosomal protein S6 kinase, 90kD, polypeptide 1	NM_002953	1.49
phosphorylase kinase, gamma 2 (testis)	NM_000294	1.49
neuronal PAS domain protein 2	NM_002518	1.49
pumilio homolog 1 (Drosophila)	BE670915	1.49
BCL2/adenovirus E1B 19kD interacting protein 1	NM_013978	1.49
proteasome (prosome, macropain) 26S subunit, ATPase, 2	NM_002803	1.49
SET translocation (myeloid leukemia-associated)	NM_003011	1.49
dynein light chain-A	NM_016141	1.49
Kaiso	BG391005	1.49
casein kinase 2, alpha 1 polypeptide	NM_001895	1.49
Ca2+-promoted Ras inactivator	NM_006989	1.49
hypothetical protein FLJ14547	AL049319	1.49
hypothetical protein MGC2668	NM_030914	1.49
pinin, desmosome associated protein	AW152664	1.49
protein phosphatase 2A, regulatory subunit B' (PR 53)	X86428	1.49
KIAA0852 protein	NM_014941	1.49
methylthioadenosine phosphorylase	AF109294	1.49
hypothetical protein FLJ13909	NM_025108	1.49
RAB6A, member RAS oncogene family	NM_002869	1.49
collagen, type IV, alpha 3 (Goodpasture antigen) binding protein	NM_005713	1.49
Alg5, S. cerevisiae, homolog of	NM_013338	1.49
benzodiazapine receptor (peripheral)	NM_000714	1.49
zinc finger protein, subfamily 1A, 1 (Ikaros)	S80876	1.49
diaphanous homolog 1 (Drosophila)	AU158818	1.49
ADP-ribosyltransferase (NAD+; poly(ADP-ribose) polymerase)-like 2	AI236912	1.49
Consensus includes gb:AA001552 /FEA=EST /DB_XREF=gi:1437017 /DB_XREF=est:ze46c08.s1 /CLONE=IMAGE:362030 /UG=Hs.59839 ESTs, Weakly similar to CGHU7L collagen alpha 1 (II	AA001552	1.49
dihydropyrimidinase-like 4	AW090187	1.49
regulator of G-protein signalling 16	U94829	1.49
Consensus includes gb:M74089.1 /DEF=Human TB1 gene mRNA, 3 end. /FEA=mRNA /GEN=TB1 /DB_XREF=gi:182400 /UG=Hs.75639 Human TB1 gene mRNA, 3 end	M74089	1.49
sorting nexin 17	NM_014748	1.49
KIAA0683 gene product	AB014583	1.49
KIAA0258 gene product	NM_014785	1.49
aminolevulinate, delta-, synthase 1	NM_000688	1.48
hypothetical protein FLJ10509	NM_018119	1.48
brefeldin A-inhibited guanine nucleotide-exchange protein 2	AV657604	1.48
inosine triphosphatase (nucleoside triphosphate pyrophosphatase)	AF219116	1.48
ribosomal protein S4, X-linked	AL035603	1.48
cyclin-dependent kinase (CDC2-like) 10	AF153430	1.48
Consensus includes gb:BG170659 /FEA=EST /DB_XREF=gi:12677362 /DB_XREF=est:602323531F1 /CLONE=IMAGE:4426926 /UG=Hs.226138 Homo sapiens mRNA; cDNA DKFZp566H244	BG170659	1.48
cytochrome c oxidase subunit VIIa polypeptide 2 like	NM_004718	1.48
hypothetical protein FLJ23045	NM_024704	1.48
nuclear transcription factor, X-box binding 1	NM_002504	1.48
ribosomal protein L7	BG389744	1.48
serologically defined colon cancer antigen 8	U79271	1.48
integrin, alpha 4 (antigen CD49D, alpha 4 subunit of VLA-4 receptor)	BG532690	1.48
hypothetical protein	NM_016618	1.48
binder of Arl Two	BF244411	1.48
oxysterol binding protein	AF185696	1.48
eukaryotic translation initiation factor 3, subunit 10 (theta, 150/170kD)	BE614908	1.48
nudix (nucleoside diphosphate linked moiety X)-type motif 9	NM_024047	1.48
chromobox homolog 4 (Pc class homolog, Drosophila)	NM_003655	1.48
protection of telomeres 1	BC002923	1.48
protein kinase, AMP-activated, beta 1 non-catalytic subunit	NM_006253	1.48
serologically defined breast cancer antigen 84	AF091085	1.48
FUS interacting protein (serine-arginine rich) 1	NM_006625	1.48
E2F transcription factor 1	NM_005225	1.48

centromere protein F (350/400kD, mitosin)	U30872	1.48
ubiquitination factor E4B (UFD2 homolog, yeast)	AW241715	1.48
N-acylsphingosine amidohydrolase (acid ceramidase)-like	AK024677	1.48
myosin, light polypeptide, regulatory, non-sarcomeric (20kD)	NM_006471	1.48
ATPase, Na+/K+ transporting, alpha 1 polypeptide	NM_000701	1.48
DKFZP564K202 protein	AL049944	1.48
KIAA0697 protein	AI986295	1.48
diacylglycerol kinase, zeta (104kD)	NM_003646	1.48
hypothetical protein DKFZp56400523	AL136619	1.48
antigen identified by monoclonal antibody Ki-67	AU147044	1.48
hypothetical protein FLJ13441	NM_023924	1.48
two-pore channel 1, homolog	NM_017901	1.48
Consensus includes gb:X69637.1 /DEF=H.sapiens mRNA sequence (16p11.2). /FEA=mRNA /DB_XREF=gi:29573 /UG=Hs.307208 H.sapiens mRNA sequence (16p11.2)	X69637	1.48
UBX domain-containing 1	NM_025241	1.48
methyl-CpG binding domain protein 1	NM_015845	1.48
KIAA0084 protein	D42043	1.48
putative DNA binding protein	AI566096	1.48
hypothetical protein FLJ13117	NM_023071	1.48
DiGeorge syndrome critical region gene D6S1; likely ortholog of mouse expressed sequence 2 embryonic lethal	NM_022719	1.48
vesicle trafficking, beach and anchor containing	W60686	1.48
holocytochrome c synthase (cytochrome c heme-lyase)	NM_005333	1.48
hypothetical protein FLJ10540	NM_018131	1.48
ornithine decarboxylase antizyme inhibitor	AA047234	1.48
eukaryotic translation initiation factor 3, subunit 7 (zeta, 66/67kD)	NM_003753	1.48
heat shock 60kD protein 1 (chaperonin)	NM_002156	1.47
CDC-like kinase 2	NM_003993	1.47
metastasis-associated 1-like 1	AA478965	1.47
hypothetical protein FLJ10535	NM_018129	1.47
adaptor-related protein complex 4, sigma 1 subunit	AF155159	1.47
HIV-1 Tat interactive protein, 60 kDa	AA886971	1.47
desmin	NM_001927	1.47
prp28, U5 snRNP 100 kd protein	NM_004818	1.47
talin-interacting protein	AI220627	1.47
F-box only protein 3	NM_012175	1.47
Consensus includes gb:AL049987.1 /DEF=Homo sapiens mRNA; cDNA DKFZp564F112 (from clone DKFZp564F112). /FEA=mRNA /DB_XREF=gi:4884238 /UG=Hs.166361 Homo sapiens mF	AL049987	1.47
hypothetical protein FLJ10111	NM_017999	1.47
thyroid hormone receptor coactivating protein	NM_006696	1.47
PTD016 protein	NM_016125	1.47
Consensus includes gb:AL021395 /DEF=Human DNA sequence from clone RP1-269M15 on chromosome 20q12-13.12 Contains a gene similar to peptidylprolyl isomerase (cyclophilin), part	AL021395	1.47
ilvB (bacterial acetolactate synthase)-like	NM_006844	1.47
Consensus includes gb:L43577 /FEA=EST /DB_XREF=gi:899066 /DB_XREF=est:L43577 /CLONE=EST500 /UG=Hs.82171 Homo sapiens (clone 115392) mRNA	L43577	1.47
cell division cycle 25C	NM_001790	1.47
RAP1, GTP-GDP dissociation stimulator 1	BC001851	1.47
ethanolamine kinase	AL137750	1.47
small acidic protein	NM_014267	1.47
CCR4-NOT transcription complex, subunit 3	AF180474	1.47
bromodomain adjacent to zinc finger domain, 1B	AI826454	1.47
basic transcription factor 3	X74070	1.47
manic fringe homolog (Drosophila)	NM_002405	1.47
microfibrillar-associated protein 1	NM_005926	1.47
phospholipid transfer protein	NM_006227	1.47
hypothetical protein FLJ21799	NM_024946	1.47
purinergic receptor P2Y, G-protein coupled, 11	NM_002566	1.47
macrophage myristoylated alanine-rich C kinase substrate	NM_023009	1.47
X-ray repair complementing defective repair in Chinese hamster cells 3	AK022829	1.47
glioma tumor suppressor candidate region gene 2	NM_015710	1.47
hypothetical protein MGC2875	NM_024073	1.47
nuclease sensitive element binding protein 1	BC002411	1.47
Friedreich ataxia	NM_000144	1.47
apolipoprotein M	AF161454	1.47
Consensus includes gb:AK001816.1 /DEF=Homo sapiens cDNA FLJ10954 fis, clone PLACE1000383, highly similar to Homo sapiens mRNA for MTMR1 protein. /FEA=mRNA /DB_XREF=gi:70	AK001816	1.47
protein phosphatase 2, regulatory subunit B (B56), epsilon isoform	NM_006246	1.47
serine/threonine kinase 19	L26260	1.47

TNF receptor-associated factor 2	NM_021138	1.47
membrane-associated tyrosine- and threonine-specific cdc2-inhibitory kinase	NM_004203	1.47
Consensus includes gb:AI907083 /FEA=EST /DB_XREF=gi:6497611 /DB_XREF=est:PM-BT134-050499-650 /UG=Hs.124620 ESTs	AI907083	1.47
zinc finger protein 131 (clone pHZ-10)	AW968301	1.47
trinucleotide repeat containing 5	BC004423	1.47
sympletin; Huntingtin interacting protein I	NM_004819	1.47
interleukin enhancer binding factor 1	NM_004514	1.47
CGI-99 protein	NM_016039	1.47
hypothetical protein FLJ13220	NM_021927	1.47
Rho-associated, coiled-coil containing protein kinase 2	D87931	1.47
phosphoinositide-3-kinase, regulatory subunit, polypeptide 1 (p85 alpha)	AI680192	1.47
ribonuclease/angiogenin inhibitor	NM_002939	1.47
Consensus includes gb:AL049761 /DEF=Human DNA sequence from clone RP5-863C7 on chromosome 20p12.3-13. Contains the CSNK2A1 gene for casein kinase 2 alpha 1 polypeptide (EC	AL049761	1.47
fatty-acid-Coenzyme A ligase, very long-chain 1	NM_003645	1.47
IGF-II mRNA-binding protein 3	NM_006547	1.46
COP9 (constitutive photomorphogenic, Arabidopsis, homolog) subunit 7B	NM_022730	1.46
DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 11 (CHL1-like helicase homolog, S. cerevisiae)	NM_030653	1.46
hypothetical protein DKFZp564D0478	AL136683	1.46
nuclear transcription factor Y, alpha	AL031778	1.46
oxidation resistance 1	NM_018002	1.46
heat shock 90kD protein 1, alpha	AF028832	1.46
metallothionein 1H	AF333388	1.46
CDC14 cell division cycle 14 homolog A (S. cerevisiae)	AF064103	1.46
plasminogen	M74220	1.46
heme-regulated initiation factor 2-alpha kinase	NM_014413	1.46
hypothetical protein, clone Telethon (Italy_841)_Strait02270_FL142	NM_020247	1.46
cutaneous T-cell lymphoma-associated tumor antigen se20-4; differentially expressed nucleolar TGF-beta1 target protein (DENTT)	NM_022117	1.46
adenosine A2a receptor	NM_000675	1.46
golgi associated, gamma adaptin ear containing, ARF binding protein 3	AW008018	1.46
DKFZP434F195 protein	BE222901	1.46
ribosomal protein L24	AI560573	1.46
actin, gamma 1	BG026805	1.46
MAX protein	M64240	1.46
DnaJ (Hsp40) homolog, subfamily B, member 6	AF080569	1.46
misato	NM_018116	1.46
mitochondrial ribosomal protein L49	NM_004927	1.46
Cluster Incl. AA524093:ng33f04.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-936607 /clone_end=3 /gb=AA524093 /gi=2265021 /ug=Hs.23158 /len=718	AA524093	1.46
hypothetical protein FLJ22246	NM_025232	1.46
prolylcarboxypeptidase (angiotensinase C)	NM_005040	1.46
WW domain binding protein 2	U79458	1.46
hypothetical protein FLJ10659	NM_018171	1.46
hypothetical protein FLJ13949	NM_025077	1.46
KIAA0429 gene product	AF116674	1.46
spectrin, alpha, non-erythrocytic 1 (alpha-fodrin)	AL110273	1.46
hypothetical protein MGC4172	NM_024308	1.46
aquaporin 3	NM_004925	1.46
Consensus includes gb:W95043 /FEA=EST /DB_XREF=gi:1424182 /DB_XREF=est:zh46c11.s1 /CLONE=IMAGE:415124 /UG=Hs.79732 fibulin 1	W95043	1.46
zinc finger protein 278	AF242522	1.46
eukaryotic translation initiation factor 3, subunit 1 (alpha, 35kD)	BC002719	1.46
hypothetical protein dJ122O8.2	NM_020466	1.45
tyrosine 3-monooxygenase/tryptophan 5-monooxygenase activation protein, theta polypeptide	AA854017	1.45
metastasis associated 1	BE561596	1.45
Consensus includes gb:AL024509 /DEF=Human DNA sequence from clone 522P13 on chromosome 6p21.31-22.3. Contains a 60S Ribosomal Protein L21 pseudogene and an HNRNP A3 (He	AL024509	1.45
hypothetical protein FLJ14972	AA535065	1.45
oxidative-stress responsive 1	NM_005109	1.45
melanoma cell adhesion molecule	BC006329	1.45
YME1-like 1 (S. cerevisiae)	AF070656	1.45
hypothetical protein FLJ10450	NM_018095	1.45
origin recognition complex, subunit 1-like (yeast)	NM_004153	1.45
paternally expressed 10	AL582836	1.45
core promoter element binding protein	AB017493	1.45
hypothetical protein FLJ12785	NM_024855	1.45
KIAA0063 gene product	NM_014876	1.45

carnitine palmitoyltransferase I, muscle	U62733	1.45
cyclin F	NM_001761	1.45
latent transforming growth factor beta binding protein 4	NM_003573	1.45
glutathione transferase zeta 1 (maleylacetoacetate isomerase)	BC001453	1.45
low density lipoprotein-related protein-associated protein 1 (alpha-2-macroglobulin receptor-associated protein 1)	NM_002337	1.45
KIAA0776 protein	AW298092	1.45
thioredoxin, mitochondrial	AF276920	1.45
casein kinase 1, alpha 1	BF341845	1.45
gb:BC002878.1 /DEF=Homo sapiens, Similar to cofactor required for Sp1 transcriptional activation, subunit 8 (34kD), clone MGC:11274, mRNA, complete cds. /FEA=mRNA /PROD=Similar t	BC002878	1.45
TRIAD3 protein	NM_019011	1.45
hypothetical protein FLJ22127	NM_022775	1.45
hypothetical protein	BC005883	1.45
PR domain containing 2, with ZNF domain	AV724783	1.45
pro-oncosis receptor inducing membrane injury gene	BG538627	1.45
hypothetical protein FLJ10547	AL049795	1.45
aquaporin 7	NM_001170	1.45
cell division cycle 2-like 5 (cholinesterase-related cell division controller)	NM_003718	1.45
RAD50 homolog (S. cerevisiae)	U63139	1.45
sialyltransferase 1 (beta-galactoside alpha-2,6-sialytransferase)	AI743792	1.45
zinc finger protein 22 (KOX 15)	NM_006963	1.45
Consensus includes gb:AA643304 /FEA=EST /DB_XREF=gi:2568522 /DB_XREF=est:nr59a09.s1 /CLONE=IMAGE:1172248 /UG=Hs.170218 KIAA0251 protein	AA643304	1.45
hypothetical protein FLJ13390 similar to PAF53	NM_022490	1.45
phosphoribosylglycinamide formyltransferase, phosphoribosylglycinamide synthetase, phosphoribosylaminoimidazole synthetase	BE966876	1.45
DNA segment, single copy, probe pH4 (transforming sequence, thyroid-1,	NM_005436	1.45
GTPase GES; REM protein	AF152863	1.45
UDP-Gal:betaGlcNAc beta 1,4- galactosyltransferase, polypeptide 1	D29805	1.45
ornithine decarboxylase antizyme 1	AF090094	1.45
Consensus includes gb:AI348006 /FEA=EST /DB_XREF=gi:4085212 /DB_XREF=est:qp61b05.s1 /CLONE=IMAGE:1927473 /UG=Hs.469 succinate dehydrogenase complex, subunit A, flavopr	AI348006	1.45
cell division cycle 25B	NM_021873	1.45
solute carrier family 25 (mitochondrial carrier; adenine nucleotide translocator), member 5	NM_001152	1.45
nuclear VCP-like	NM_002533	1.45
RPB5-mediating protein	AW514900	1.45
KIAA0372 gene product	BE566023	1.45
carboxypeptidase D	NM_001304	1.45
hypothetical protein	BE732345	1.44
hypothetical protein FLJ13044	NM_024698	1.44
clone FLB5214	NM_016418	1.44
Consensus includes gb:AA527499 /FEA=EST /DB_XREF=gi:2269568 /DB_XREF=est:ng41f07.s1 /CLONE=IMAGE:937381 /UG=Hs.301667 Homo sapiens mRNA; cDNA DKFZp566l043 (from	AA527499	1.44
plakophilin 4	NM_003628	1.44
RAN binding protein 16	H38643	1.44
adaptor-related protein complex 3, beta 1 subunit	NM_003664	1.44
major histocompatibility complex, class I, F	AI669379	1.44
hypothetical protein MGC4825	AF061264	1.44
ras homolog gene family, member F (in filopodia)	NM_019034	1.44
HSPC022 protein	BE138888	1.44
general transcription factor IIIC, polypeptide 1 (alpha subunit, 220kD)	NM_001520	1.44
KIAA0171 gene product	BC004467	1.44
solute carrier family 2 (facilitated glucose transporter), member 1	NM_006516	1.44
myelin basic protein	AW070431	1.44
CGI-141 protein	NM_016072	1.44
putative dimethyladenosine transferase	BC002841	1.44
HGF activator	NM_001528	1.44
hypothetical protein FLJ20374	NM_017793	1.44
Consensus includes gb:AK024269.1 /DEF=Homo sapiens cDNA FLJ14207 fis, clone NT2RP3003185, weakly similar to TROPOMYOSIN 1, FUSION PROTEIN 33. /FEA=mRNA /DB_XREF=gi:104	AK024269	1.44
hypothetical protein DKFZp434E2216	BC001043	1.44
RAD23 homolog B (S. cerevisiae)	NM_002874	1.44
enolase 1, (alpha)	BC005884	1.44
eukaryotic translation initiation factor 4E-like 3	BF516289	1.44
ATP-binding cassette, sub-family B (MDR/TAP), member 7	AB005289	1.44
aspartyl aminopeptidase	AF005050	1.44
zinc finger protein 273	AU132789	1.44
LAG1 longevity assurance homolog 2 (S. cerevisiae)	AK001105	1.44
SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily e, member 1	NM_003079	1.44

ceroid-lipofuscinosis, neuronal 3, juvenile (Batten, Spielmeyer-Vogt disease)	AF077973	1.44
interferon-stimulated protein, 15 kDa	NM_005101	1.44
KIAA0678 protein	AB014578	1.44
CGI-40 protein	AA150165	1.44
KIAA0923 protein	AU152583	1.44
BCL2-associated X protein	NM_004324	1.44
hypothetical protein FLJ20850	AI744084	1.44
putative nucleic acid binding protein RY-1	BG252325	1.44
UDP-glucose ceramide glucosyltransferase	NM_003358	1.44
heparan sulfate 2-O-sulfotransferase	AW151887	1.44
folylpolyglutamate synthase	NM_004957	1.44
mucosa associated lymphoid tissue lymphoma translocation gene 1	NM_006785	1.44
developmentally regulated GTP binding protein 2	BF223206	1.44
Consensus includes gb:AL133053.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434H1215 (from clone DKFZp434H1215); partial cds. /FEA=mRNA /GEN=DKFZp434H1215 /PROD=hypothetic	AL133053	1.44
RAD1 homolog (S. pombe)	AF074717	1.43
cAMP responsive element binding protein 3 (human)	AF029674	1.43
proline arginine-rich and leucine-rich repeat protein	U41344	1.43
sulfotransferase family, cytosolic, 1A, phenol-preferring, member 3	U08032	1.43
KIAA0630 protein	AI393355	1.43
hypothetical protein FLJ13910	BF671894	1.43
colony stimulating factor 2 receptor, beta, low-affinity (granulocyte-macrophage)	AV756141	1.43
nuclear transcription factor Y, gamma	U78774	1.43
nudix (nucleoside diphosphate linked moiety X)-type motif 4	AF191653	1.43
proteasome (prosome, macropain) subunit, alpha type, 3	NM_002788	1.43
succinate dehydrogenase complex, subunit D, integral membrane protein	NM_003002	1.43
steroidogenic acute regulatory protein related	NM_006804	1.43
phosphate cytidylyltransferase 2, ethanolamine	BC000351	1.43
cell division cycle 42 (GTP binding protein, 25kD)	BC002711	1.43
high-mobility group (nonhistone chromosomal) protein 14	NM_004965	1.43
Consensus includes gb:AI858004 /FEA=EST /DB_XREF=gi:5511620 /DB_XREF=est:wj69b09.x1 /CLONE=IMAGE:2408057 /UG=Hs.284277 Homo sapiens immunoglobulin mu chain antibod	AI858004	1.43
zinc finger protein 103 homolog (mouse)	NM_005667	1.43
aprataxin	NM_017692	1.43
leucyl-tRNA synthetase, mitochondrial	D21851	1.43
DKFZP727M231 protein	AL117480	1.43
HIV-1 Tat interactive protein 2, 30 kDa	NM_006410	1.43
cullin 4A	AL037208	1.43
ARP1 actin-related protein 1 homolog A, centractin alpha (yeast)	NM_005736	1.43
HSPC166 protein	NM_014186	1.43
BTB (POZ) domain containing 2	NM_017797	1.43
hypothetical protein FLJ14393	BE966695	1.43
hypothetical protein FLJ11273	NM_018374	1.43
SAR1 protein	NM_020150	1.43
hypothetical protein FLJ11126	BC005162	1.43
inositol 1,4,5-triphosphate receptor, type 3	D26351	1.43
splicing factor, arginine/serine-rich 8 (suppressor-of-white-apricot homolog, Drosophila)	NM_004592	1.43
protein kinase C, alpha	NM_002737	1.43
lymphoid enhancer-binding factor 1	AF294627	1.43
nuclear autoantigenic sperm protein (histone-binding)	AW003362	1.43
glutamine-fructose-6-phosphate transaminase 1	NM_002056	1.43
hypothetical protein FLJ14827	AI937468	1.43
KIAA0846 protein	NM_015376	1.43
ubiquitously-expressed transcript	NM_004182	1.43
proliferating cell nuclear antigen	NM_002592	1.43
runt-related transcription factor 3	NM_004350	1.43
fragile X mental retardation 1	AI743037	1.43
KIAA1253 protein	AF164794	1.43
hydroxymethylbilane synthase	NM_000190	1.43
TS4 protein	NM_015698	1.43
RNA polymerase I transcription factor RRN3	AF001549	1.43
hypothetical protein FLJ10081	NM_017991	1.43
keratin 6B	L42612	1.43
integrin, beta 1 (fibronectin receptor, beta polypeptide, antigen CD29 includes MDF2, MSK12)	BG334196	1.43
N-acetylneuraminic acid phosphate synthase; sialic acid synthase	NM_018946	1.43

COP9 (constitutive photomorphogenic, Arabidopsis, homolog) subunit 5	NM_006837	1.43
fibroblast growth factor receptor 2 (bacteria-expressed kinase, keratinocyte growth factor receptor, craniofacial dysostosis 1, Crouzon syndrome, Pfeiffer syndrome, Jackson-Weiss syndrome	NM_022975	1.43
zinc finger protein 151 (pHZ-67)	NM_003443	1.42
putative mitochondrial outer membrane protein import receptor	BG469030	1.42
secretory carrier membrane protein 1	AF005037	1.42
MADS box transcription enhancer factor 2, polypeptide C (myocyte enhancer factor 2C)	AL536517	1.42
phosphatidylinositol glycan, class N	NM_012327	1.42
Spi-B transcription factor (Spi-1/PU.1 related)	NM_003121	1.42
RNA binding motif protein 9	AW149379	1.42
fibroblast growth factor receptor 2 (bacteria-expressed kinase, keratinocyte growth factor receptor, craniofacial dysostosis 1, Crouzon syndrome, Pfeiffer syndrome, Jackson-Weiss syndrome	AB030077	1.42
ADP-ribosyltransferase (NAD+; poly (ADP-ribose) polymerase)-like 2	NM_005484	1.42
chemokine-like factor 1	NM_016326	1.42
programmed cell death 6 interacting protein	NM_013374	1.42
F-box only protein 9	AF176704	1.42
hypothetical protein PP591	NM_025207	1.42
karyopherin (importin) beta 3	AF339834	1.42
heterogeneous nuclear ribonucleoprotein A/B	NM_004499	1.42
extra spindle poles, S. cerevisiae, homolog of	D79987	1.42
cullin 2	NM_003591	1.42
SET translocation (myeloid leukemia-associated)	D45198	1.42
minor histocompatibility antigen HA-1	BE349017	1.42
zinc finger protein 79 (pT7)	X65232	1.42
carboxypeptidase D	AA897514	1.42
potassium inwardly-rectifying channel, subfamily J, member 4	Z97056	1.42
host cell factor C1 (VP16-accessory protein)	NM_005334	1.42
Consensus includes gb:AI052536 /FEA=EST /DB_XREF=gi:3308527 /DB_XREF=est:oz27g09.x1 /CLONE=IMAGE:1676608 /UG=Hs.31834 Homo sapiens clone 25129 mRNA sequence	AI052536	1.42
KIAA0415 gene product	AB007875	1.42
RAB48, member RAS oncogene family	NM_016154	1.42
hypothetical protein FLJ21613 similar to rat corneal wound healing related protein	NM_021929	1.42
cyclin-dependent kinase 4	U81031	1.42
solute carrier family 25 (mitochondrial carrier; oxoglutarate carrier), member 11	AF070548	1.42
suppressor of Ty 6 homolog (S. cerevisiae)	D79984	1.42
ring finger protein 13	AF070558	1.42
Consensus includes gb:AL050217.1 /DEF=Homo sapiens mRNA; cDNA DKFZp586I0523 (from clone DKFZp586I0523). /FEA=mRNA /DB_XREF=gi:4884458 /UG=Hs.308467 Homo sapiens n	AL050217	1.42
ATP synthase, H+ transporting, mitochondrial F0 complex, subunit b, isoform 1	BC005960	1.42
cell division cycle 25A	AI343459	1.42
KIAA0560 gene product	BG260519	1.42
MADS box transcription enhancer factor 2, polypeptide A (myocyte enhancer factor 2A)	X63381	1.42
Huntingtin interacting protein K	NM_016400	1.42
CGI-112 protein	NM_016049	1.42
trefoil factor 1 (breast cancer, estrogen-inducible sequence expressed in)	NM_003225	1.42
testis enhanced gene transcript (BAX inhibitor 1)	AF033095	1.42
sterol regulatory element binding transcription factor 1	NM_004176	1.42
adaptor-related protein complex 3, beta 1 subunit	AW058575	1.42
TRAM-like protein	NM_012288	1.42
pre-mRNA processing factor 31 homolog (yeast)	NM_015629	1.42
B cell RAG associated protein	NM_014863	1.42
hypothetical protein FLJ21952	NM_022494	1.42
Consensus includes gb:L43578.1 /DEF=Homo sapiens (clone 115392) mRNA. /FEA=mRNA /DB_XREF=gi:899067 /UG=Hs.82171 Homo sapiens (clone 115392) mRNA	L43578	1.42
KIAA0854 protein	NM_014943	1.42
golgi autoantigen, golgin subfamily a, 5	NM_005113	1.42
hypothetical protein DKFZp586F2423	BE675139	1.42
Consensus includes gb:AI076186 /FEA=EST /DB_XREF=gi:3405364 /DB_XREF=est:oz01g01.x1 /CLONE=IMAGE:1674096 /UG=Hs.284275 Homo sapiens PAK2 mRNA, complete cds /FL=gt	AI076186	1.42
hypothetical protein FLJ14007	NM_024699	1.42
KIAA0211 gene product	BF508616	1.42
hypothetical protein FLJ21459	NM_024521	1.42
stannin	AF070673	1.42
hypothetical protein FLJ12587	AL136796	1.42
zinc finger protein with interaction domain	NM_006626	1.42
hypothetical protein FLJ20189	NM_017704	1.42
hypothetical protein FLJ20445	NM_017824	1.42
KIAA0102 gene product	NM_014752	1.42
peroxisomal acyl-CoA thioesterase 1	NM_005469	1.42

spermine synthase	NM_004595	1.42
synaptobrevin-like 1	NM_005638	1.42
RNA binding motif, single stranded interacting protein 2	D28483	1.42
hypothetical protein FLJ10704	NM_018185	1.42
survival of motor neuron 1, telomeric	NM_000344	1.42
metallothionein 1F (functional)	M10943	1.42
hypothetical protein FLJ22531	NM_024650	1.41
heat shock 105kD	NM_006644	1.41
retinoblastoma-like 2 (p130)	BF110947	1.41
aspartate beta-hydroxylase	AF289489	1.41
H2A histone family, member Z	BF718636	1.41
hypothetical protein FLJ21213	NM_022900	1.41
BCL2-antagonist of cell death	U66879	1.41
KIAA0252 protein	AI192081	1.41
crystallin, alpha A	AP001748	1.41
transcription factor 8 (represses interleukin 2 expression)	U12170	1.41
Ras homolog enriched in brain 2	BF593727	1.41
ring finger protein 24	NM_007219	1.41
G protein-coupled receptor 65	NM_003608	1.41
replication factor C (activator 1) 5 (36.5kD)	BC001866	1.41
FAST kinase	BC000377	1.41
hypothetical protein FLJ20411	AB050442	1.41
SH3-domain, GRB2-like, endophilin B1	AF257318	1.41
leucyl/cystinyl aminopeptidase	NM_005575	1.41
amplified in osteosarcoma	NM_006812	1.41
cyclin G associated kinase	D88435	1.41
hypothetical protein	BG223334	1.41
gb:NM_015930.1 /DEF=Homo sapiens vanilloid receptor-like protein 1 (VRL-1), mRNA /FEA=mRNA /GEN=VRL-1 /PROD=vanilloid receptor-like protein 1 /DB_XREF=gi:7706764 /UG=Hs.2	NM_015930	1.41
exosome component Rrp41	AL039469	1.41
MLL septin-like fusion	AB023208	1.41
leucine zipper transcription factor-like 1	NM_020347	1.41
mitogen-activated protein kinase-activated protein kinase 2	AI141802	1.41
hypothetical protein 38408_6	BC001298	1.41
WNT1 inducible signaling pathway protein 2	NM_003881	1.41
actin related protein 2/3 complex, subunit 4 (20 kD)	NM_005718	1.41
zinc finger protein 143 (clone pHZ-1)	AW162015	1.41
high-mobility group (nonhistone chromosomal) protein 1	AC007277	1.41
mutS homolog 6 (E. coli)	NM_000179	1.41
hypothetical protein FLJ20580	AW157077	1.41
adenomatosis polyposis coli	M74088	1.41
ornithine decarboxylase antizyme inhibitor	NM_015878	1.41
HCF-binding transcription factor Zhangfei	AW204564	1.41
N-ethylmaleimide-sensitive factor attachment protein, alpha	NM_003827	1.41
8-oxoguanine DNA glycosylase	NM_016821	1.41
protein tyrosine phosphatase, non-receptor type 11	L07527	1.41
hypothetical protein FLJ22408	NM_024794	1.41
caspase 6, apoptosis-related cysteine protease	BC000305	1.41
Rab geranylgeranyltransferase, beta subunit	U49245	1.41
Consensus includes gb:AA906578 /FEA=EST /DB_XREF=gi:3042164 /DB_XREF=est:ol22c06.s1 /CLONE=IMAGE:1524202 /UG=Hs.23158 ESTs	AA906578	1.41
acyl-Coenzyme A dehydrogenase, very long chain	NM_000018	1.41
NADH dehydrogenase (ubiquinone) Fe-S protein 4 (18kD) (NADH-coenzyme Q reductase)	BC005270	1.41
hypothetical protein MGC2747	NM_024104	1.41
Mouse Mammary Tumor Virus Receptor homolog	BF974389	1.41
cadherin 6, type 2, K-cadherin (fetal kidney)	BC000019	1.41
HSPC035 protein	NM_016127	1.41
nuclear receptor subfamily 2, group F, member 6	BC002669	1.41
v-crk avian sarcoma virus CT10 oncogene homolog	BF304695	1.40
golgi associated, gamma adaptin ear containing, ARF binding protein 2	AF190863	1.40
torsin family 1, member B (torsin B)	AF317129	1.40
capping protein (actin filament) muscle Z-line, beta	AL572341	1.40
hypothetical protein FLJ10539	NM_018130	1.40
hypothetical protein FLJ10402	NM_018085	1.40
meningioma expressed antigen 6 (coiled-coil proline-rich)	U73682	1.40

proprotein convertase subtilisin/kexin type 1 inhibitor	NM_013271	1.40
regulator of G-protein signalling 10	NM_002925	1.40
KIAA0978 protein	AL117518	1.40
HIV TAT specific factor 1	AI373539	1.40
ribosomal protein S6 kinase, 70kD, polypeptide 1	M60725	1.40
KIAA0728 protein	BC004912	1.40
guanine nucleotide binding protein (G protein), beta polypeptide 1	NM_002074	1.40
RAP2A, member of RAS oncogene family	BE669921	1.40
BCL2-associated athanogene 2	AF095192	1.40
nuclear factor of kappa light polypeptide gene enhancer in B-cells inhibitor, epsilon	NM_004556	1.40
cell division cycle 27	AL566824	1.40
WD repeat domain 8	NM_017818	1.40
CG6 triplet repeat binding protein 1	NM_003663	1.40
replication factor C (activator 1) 1 (145kD)	L14922	1.40
v-ral simian leukemia viral oncogene homolog B (ras related; GTP binding protein)	NM_002881	1.40
I-kappa-B-interacting Ras-like protein 2	AA452565	1.40
CD72 antigen	AF283777	1.40
NAD kinase	AI239732	1.40
SET domain, bifurcated 1	NM_012432	1.40
regulator of G-protein signalling 2, 24kD	NM_002923	1.40
uncharacterized bone marrow protein BM045 .	NM_018459	1.40
Consensus includes gb:AL042729 /FEA=EST /DB_XREF=gi:5935545 /DB_XREF=est:DKFZp43480222_s1 /CLONE=DKFZp43480222 /UG=Hs.112023 ESTs	AL042729	1.40
KIAA1033 protein	AL137753	1.40
PAI-1 mRNA-binding protein	AF151813	1.40
tubulin, beta, 2	BC004188	1.40
vacuolar protein sorting 35 (yeast)	NM_018206	1.40
retinitis pigmentosa 2 (X-linked recessive)	NM_006915	1.40
proteasome (prosome, macropain) 26S subunit, non-ATPase, 12	AI446530	1.40
chloride channel 3	NM_001829	1.40
peptidylprolyl isomerase E (cyclophilin E)	NM_006112	1.40
transcriptional co-activator with PDZ-binding motif (TAZ)	BF674349	1.40
Consensus includes gb:U21915.1 /DEF=Human chromosome 5q13.1 clone 5G8 mRNA. /FEA=mRNA /DB_XREF=gi:736412 /UG=Hs.14658 Human chromosome 5q13.1 clone 5G8 mRNA	U21915	1.40
hypoxia-inducible factor 1, alpha subunit inhibitor	W37897	1.40
poly(A) binding protein, cytoplasmic, pseudogene 3	U64661	1.40
degenerative spermatocyte homolog, lipid desaturase (Drosophila)	NM_003676	1.40
jumonji homolog (mouse)	NM_004973	1.40
hypothetical protein FLJ23338	NM_024658	1.40
Rab geranylgeranyltransferase, beta subunit	U49245	1.40
hypothetical protein FLJ13433	NM_022496	1.40
serine (or cysteine) proteinase inhibitor, clade B (ovalbumin), member 1	AI554300	1.40
IDN3 protein	AW339587	1.40
CGI-105 protein	NM_016044	1.40
zinc finger protein 215	NM_013250	1.40
sentrin/SUMO-specific protease 3	NM_015670	1.40
hypothetical protein MGC4606	NM_024516	1.40
HEMK homolog 7kb	W93807	1.40
thymopoietin	BG391171	1.40
putative nucleic acid binding protein RY-1	X76302	1.40
thioredoxin interacting protein	AA812232	1.40
H2B histone family, member R	NM_021058	1.39
splicing factor, arginine/serine-rich 2	BG254869	1.39
phosphatidylinositol binding clathrin assembly protein	AV722190	1.39
HSPC125 protein	NM_014165	1.39
nucleolar protein GU2	AF334103	1.39
hypothetical protein FLJ12666	NM_024595	1.39
KIAA0652 gene product	NM_014741	1.39
v-ets erythroblastosis virus E26 oncogene homolog 2 (avian)	NM_005239	1.39
UDP-galactose transporter related	NM_005827	1.39
KIAA0187 gene product	NM_014753	1.39
karyopherin (importin) beta 1	AA861608	1.39
insulin-degrading enzyme	NM_004969	1.39
ferritin, heavy polypeptide 1	AA083483	1.39
DEK oncogene (DNA binding)	NM_003472	1.39

annexin A7	J04543	1.39
elongation of very long chain fatty acids (FEN1/Elo2, SUR4/Elo3, yeast)-like 1	H93026	1.39
LIM domain kinase 1	AF134379	1.39
solute carrier family 35 (UDP-galactose transporter), member 2	D84454	1.39
Splicing factor, arginine/serine-rich, 46kD	BC006181	1.39
hypothetical protein FLJ23399	NM_022763	1.39
KIAA0212 gene product	NM_014674	1.39
N-acetylglucosaminidase, alpha- (Sanfilippo disease IIIB)	NM_000263	1.39
growth factor independent 1B (potential regulator of CDKN1A, translocated in CML)	NM_004188	1.39
Consensus includes gb:AF189251.1 /DEF=Homo sapiens cytomegalovirus partial fusion receptor mRNA, partial cds. /FEA=mRNA /PROD=cytomegalovirus partial fusion receptor /DB_XREF=	AF189251	1.39
hypothetical protein MGC2780	NM_025266	1.39
transgelin 2	BC002616	1.39
sialoadhesin	NM_023068	1.39
linker for activation of T cells	AF036905	1.39
CGI-07 protein	NM_015938	1.39
nudix (nucleoside diphosphate linked moiety X)-type motif 6	NM_007083	1.39
casein kinase 2, alpha 1 polypeptide	AI631874	1.39
hypothetical protein PRO2900	AA515560	1.39
hemochromatosis	AF149804	1.39
Consensus includes gb:AL023584 /DEF=Human DNA sequence from clone 67K17 on chromosome 6q24.1-24.3. Contains the HIVEP2 (Schnurri-2) gene for HIV type 1 Enhancer-binding Prot	AL023584	1.39
exonuclease 1	NM_003686	1.39
KIAA0033 protein	AU153138	1.39
CLIP-associating protein 2	AI741784	1.39
KIAA1096 protein	BE729523	1.39
adenylate cyclase 3	AF033861	1.39
KIAA0869 protein	AB020676	1.39
zinc finger protein 36, C3H type-like 2	AI356398	1.39
methionine-tRNA synthetase	AA621558	1.39
O-linked N-acetylglucosamine (GlcNAc) transferase (UDP-N-acetylglucosamine:polypeptide-N-acetylglucosaminyl transferase)	BF001665	1.39
ribosomal protein S23	NM_001025	1.39
hypothetical protein FLJ22693	NM_022750	1.39
protein phosphatase 3 (formerly 2B), catalytic subunit, alpha isoform (calcineurin A alpha)	AA911231	1.39
programmed cell death 10	BC002506	1.39
hypothetical protein	AA046752	1.39
ribosomal protein L32	NM_000994	1.38
hypothetical protein FLJ20657	NM_017921	1.38
APG5 autophagy 5-like (S. cerevisiae)	AK001899	1.38
ubiquitin specific protease 8	NM_005154	1.38
choline kinase	NM_001277	1.38
ceroid-lipofuscinosis, neuronal 2, late infantile (Jansky-Bielschowsky disease)	BG231932	1.38
Consensus includes gb:AK026168.1 /DEF=Homo sapiens cDNA: FLJ22515 fis, clone HRC12122, highly similar to AF052101 Homo sapiens clone 23872 mRNA sequence. /FEA=mRNA /DB_X	AK026168	1.38
CGI-49 protein	AL572542	1.38
hypothetical protein FLJ20399	AA886893	1.38
frequently rearranged in advanced T-cell lymphomas	NM_005479	1.38
hypothetical protein	AL544094	1.38
branched chain alpha-ketoacid dehydrogenase kinase	NM_005881	1.38
succinate-CoA ligase, GDP-forming, alpha subunit	NM_003849	1.38
hqp0256 protein	NM_016355	1.38
mitofusin 2	AK021947	1.38
hypothetical protein MGC4054	NM_024341	1.38
nitrogen fixation cluster-like	AY009128	1.38
KIAA0226 gene product	AI798908	1.38
adenosine deaminase, RNA-specific	NM_001111	1.38
SON DNA binding protein	NM_003103	1.38
VRK3 for vaccinia related kinase 3	BF062886	1.38
cleavage and polyadenylation specific factor 4, 30kD subunit	NM_006693	1.38
coatomer protein complex, subunit beta 2 (beta prime)	NM_004766	1.38
hypothetical protein similar to RNA-binding protein lark	BC003503	1.38
Ras-GTPase activating protein SH3 domain-binding protein 2	AU149503	1.38
phosphatase and tensin homolog (mutated in multiple advanced cancers 1)	BC005821	1.38
Consensus includes gb:AU148611 /FEA=EST /DB_XREF=gi:11010132 /DB_XREF=est:AU148611 /CLONE=NT2RM4000607 /UG=Hs.296832 Human pTR7 mRNA for repetitive sequence	AU148611	1.38
LIM domain kinase 2	AL117466	1.38
inhibitor of kappa light polypeptide gene enhancer in B-cells, kinase beta	AF080158	1.38

muscle specific gene	AF085358	1.38
LAT1-3TM protein	NM_031211	1.38
mitochondrial ribosomal protein L3	BC003375	1.38
H1 histone family, member 4	AL353759	1.38
DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 3	AF061337	1.38
hypothetical protein FLJ23109	NM_024814	1.38
polymerase (RNA) II (DNA directed) polypeptide K (7.0kD)	AL558030	1.38
Consensus includes gb:AL031602 /DEF=Human DNA sequence from clone RP5-1174N9 on chromosome 1p34.1-35.3. Contains the gene for a novel protein with IBR domain, a (pseudo?) ge	AL031602	1.38
nuclear receptor subfamily 3, group C, member 1	NM_000176	1.38
KIAA0592 protein	AL080183	1.38
postmeiotic segregation increased 2-like 11	U38980	1.38
secretory carrier membrane protein 1	AV745949	1.38
Consensus includes gb:AL080112.1 /DEF=Homo sapiens mRNA; cDNA DKFZp586H0722 (from clone DKFZp586H0722). /FEA=mRNA /DB_XREF=gi:5262539 /UG=Hs.332731 Homo sapiens	AL080112	1.38
nuclear receptor coactivator 2	NM_006540	1.38
CD1A antigen, a polypeptide	M28825	1.38
methylene tetrahydrofolate dehydrogenase (NAD+ dependent), methenyltetrahydrofolate cyclohydrolase	NM_006636	1.38
calpastatin	AF327443	1.38
hypothetical protein FLJ12528	NM_025150	1.38
KIAA0601 protein	AB011173	1.38
hypothetical protein MGC3162	NM_024078	1.38
Ewing sarcoma breakpoint region 1	AF327066	1.38
lecithin-cholesterol acyltransferase	NM_000229	1.38
protein phosphatase 2, regulatory subunit B (B56), gamma isoform	NM_002719	1.38
SH2-B homolog	AF227968	1.38
Consensus includes gb:AK022061.1 /DEF=Homo sapiens cDNA FLJ11999 fis, clone HEMBB1001527. /FEA=mRNA /DB_XREF=gi:10433381 /UG=Hs.306622 Homo sapiens cDNA FLJ11999 f	AK022061	1.38
hypothetical protein FLJ10587	NM_018149	1.38
nuclear receptor coactivator 3	AI438999	1.38
KIAA0062 protein	D31887	1.38
hypothetical protein MGC10772	NM_030567	1.38
Consensus includes gb:AC004144 /DEF=Homo sapiens chromosome 19, overlapping cosmids R28707 and R34001 /FEA=CDS /DB_XREF=gi:2887496 /UG=Hs.307350 Homo sapiens chrom	AC004144	1.37
phosphoglycerate kinase 1	NM_000291	1.37
RNA binding protein S1, serine-rich domain	NM_006711	1.37
immediate early response 5	NM_016545	1.37
PAI-1 mRNA-binding protein	AF131807	1.37
suppression of tumorigenicity 14 (colon carcinoma, matriptase, epithin)	U20428	1.37
core-binding factor, runt domain, alpha subunit 2; translocated to, 3	NM_005187	1.37
KIAA0076 gene product	NM_014780	1.37
Consensus includes gb:AI245523 /FEA=EST /DB_XREF=gi:3840920 /DB_XREF=est:qk25c04.x1 /CLONE=IMAGE:1869990 /UG=Hs.155572 Human clone 23801 mRNA sequence	AI245523	1.37
DnaJ (Hsp40) homolog, subfamily A, member 3	NM_005147	1.37
ClpX caseinolytic protease X homolog (E. coli)	NM_006660	1.37
dynamin 2	NM_004945	1.37
KIAA0794 protein	AK026451	1.37
hypothetical protein FLJ21940	NM_022828	1.37
transmembrane trafficking protein	NM_006827	1.37
hypothetical protein MGC3121	NM_024031	1.37
KIAA0852 protein	AC004542	1.37
hypothetical protein	AF217514	1.37
ATP synthase, H+ transporting, mitochondrial F1 complex, O subunit (oligomycin sensitivity conferring protein)	S77356	1.37
golgi SNAP receptor complex member 2	BE895285	1.37
G protein-coupled receptor 87	NM_023915	1.37
pituitary tumor-transforming 3	NM_021000	1.37
guanine nucleotide binding protein 10	NM_004125	1.37
KIAA0295 protein	AA913513	1.37
Consensus includes gb:AL137403.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434L092 (from clone DKFZp434L092). /FEA=mRNA /DB_XREF=gi:6807951 /UG=Hs.306457 Homo sapiens mR	AL137403	1.37
acetyl-Coenzyme A acyltransferase 2 (mitochondrial 3-oxoacyl-Coenzyme A thiolase)	NM_006111	1.37
Protein P3	NM_019848	1.37
small nuclear ribonucleoprotein polypeptide C	NM_003093	1.37
zinc finger protein 36, C3H type-like 1	BG250310	1.37
mannosyl (alpha-1,6-)-glycoprotein beta-1,2-N-acetylglucosaminyltransferase	BC006390	1.37
calpain 7	NM_014296	1.37
6.2 kd protein	NM_019059	1.37
far upstream element (FUSE) binding protein 3	U69127	1.37
retinoblastoma binding protein 8	NM_002894	1.37

small acidic protein	BE748698	1.37
E4F transcription factor 1	NM_004424	1.37
signal transducer and activator of transcription 6, interleukin-4 induced	BC004973	1.37
syntaxin 8	NM_004853	1.37
DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 11 (CHL1-like helicase homolog, S. cerevisiae)	NM_004399	1.37
Consensus includes gb:AL008627 /DEF=Human DNA sequence from PAC 13062 on chromosome 6p22.2-22.3. Contains ribosomal protein L29 pseudogene, ESTs and STSs /FEA=mRNA /DB_	AL008627	1.37
thyroid hormone receptor interactor 4	NM_016213	1.37
ribosomal protein S21	AW024383	1.37
gb:U18810.1 /DEF=Human PACAP type-3VIP type-2 receptor mRNA, complete cds. /FEA=mRNA /PROD=PACAP type-3VIP type-2 receptor /DB_XREF=gi:1684934 /FL=gb:U18810.1	U18810	1.37
MYC-associated zinc finger protein (purine-binding transcription factor)	NM_002383	1.37
hypothetical protein FLJ11240	NM_018368	1.37
potassium channel, subfamily K, member 13	NM_022054	1.37
aspartylglucosaminidase	NM_000027	1.37
KIAA0792 gene product	AW510783	1.37
c-myc binding protein	AL525412	1.37
serum response factor (c-fos serum response element-binding transcription factor)	NM_003131	1.37
formin-binding protein 3	AA675900	1.37
hypothetical protein FLJ10849	NM_018243	1.37
KIAA1052 protein	NM_014956	1.37
opioid growth factor receptor	NM_007346	1.37
hypothetical protein FLJ22294	NM_030825	1.37
SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily a, member 4	AI831675	1.37
protein phosphatase 1, regulatory (inhibitor) subunit 15A	NM_014330	1.37
ATPase, Ca++ transporting, type 2C, member 1	AF189723	1.37
Consensus includes gb:AI275383 /DEF=Homo sapiens partial IGVH3 gene for immunoglobulin heavy chain V region, case 1, cell Mo IV 72 /FEA=CDS /DB_XREF=gi:7573027 /UG=Hs.2723SI	AI275383	1.37
hypothetical protein FLJ20160	NM_017694	1.37
translocation protein 1	U93239	1.37
ceroid-lipofuscinosis, neuronal 2, late infantile (Jansky-Bielschowsky disease)	AA602532	1.37
heparan sulfate 2-O-sulfotransferase	AK023260	1.36
mitofusin 1	NM_017927	1.36
suppressor of Ty 6 homolog (S. cerevisiae)	AI494567	1.36
PDZ-binding kinase; T-cell originated protein kinase	NM_018492	1.36
KIAA0406 gene product	AB007866	1.36
sterol O-acyltransferase (acyl-Coenzyme A: cholesterol acyltransferase) 1	L21934	1.36
actin cross-linking factor	AB029290	1.36
ribosomal protein L41	NM_021104	1.36
chromosome 11 hypothetical protein ORF4	AA479495	1.36
male-enhanced antigen	NM_014623	1.36
Fas (TNFRSF6) associated factor 1	NM_007051	1.36
KIAA0365 gene product	BG252853	1.36
CUG triplet repeat, RNA binding protein 1	NM_006560	1.36
ubiquitin-conjugating enzyme E2 variant 1	AL121873	1.36
E74-like factor 4 (ets domain transcription factor)	U32645	1.36
hypothetical protein 24432	NM_022914	1.36
RNA binding protein	AF251062	1.36
heat shock 70kD protein 9B (mortalin-2)	BC000478	1.36
uncharacterized bone marrow protein BM046	NM_018460	1.36
splicing factor 3b, subunit 1, 155kD	NM_012433	1.36
carnitine palmitoyltransferase I, muscle	U62733	1.36
ribosomal protein S27a	NM_002954	1.36
membrane cofactor protein (CD46, trophoblast-lymphocyte cross-reactive antigen)	AL570661	1.36
ubiquitin-conjugating enzyme E2G 1 (UBC7 homolog, C. elegans)	BC002775	1.36
tropomodulin 3 (ubiquitous)	NM_014547	1.36
heterogeneous nuclear ribonucleoprotein H3 (2H9)	AF132362	1.36
centrosomal protein 2	AF022655	1.36
hypothetical protein FLJ10307	NM_018053	1.36
CGI-02 protein	NM_012123	1.36
3-hydroxy-3-methylglutaryl-Coenzyme A reductase	NM_000859	1.36
nuclear phosphoprotein similar to S. cerevisiae PWP1	NM_007062	1.36
exportin 1 (CRM1 homolog, yeast)	D89729	1.36
hypothetical protein FLJ21945	NM_025203	1.36
KIAA0193 gene product	NM_014766	1.36
KIAA0483 protein	NM_015176	1.36

KIAA0116 protein	AL581473	1.36
NS1-associated protein 1	AI472757	1.36
transcription factor 1, hepatic; LF-B1, hepatic nuclear factor (HNF1), albumin proximal factor	M57732	1.36
kynurenine 3-monooxygenase (kynurenine 3-hydroxylase)	AI074145	1.36
hypothetical protein FLJ10466	NM_018100	1.36
KIAA0676 protein	NM_015043	1.36
phosphomannomutase 2	NM_000303	1.36
filamin A, alpha (actin binding protein 280)	NM_001456	1.36
ret finger protein	AL523814	1.36
protein phosphatase 1, regulatory (inhibitor) subunit 2	BF966540	1.36
Consensus includes gb:X81636.1 /DEF=H.sapiens clathrin light chain a gene. /FEA=mRNA /DB_XREF=gi:704460 /UG=Hs.285688 H.sapiens clathrin light chain a gene	X81636	1.36
COP9 (constitutive photomorphogenic, Arabidopsis, homolog) subunit 4	NM_016129	1.36
melanoma antigen, family A, 12	U82671	1.36
adaptor-related protein complex 1, sigma 1 subunit	NM_001283	1.36
DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 10 (RNA helicase)	NM_004398	1.36
leukocyte immunoglobulin-like receptor, subfamily A (with TM domain), member 2	U82278	1.35
macrophage stimulating 1 (hepatocyte growth factor-like)	NM_020998	1.35
30 kDa protein	NM_018447	1.35
Sac domain-containing inositol phosphatase 2	NM_014937	1.35
Consensus includes gb:AA650558 /FEA=EST /DB_XREF=gi:2577886 /DB_XREF=est:nt02g10.s1 /CLONE=IMAGE:1192002 /UG=Hs.325202 ESTs, Highly similar to GBAS_HUMAN GUANINE N	AA650558	1.35
copine III	NM_003909	1.35
upstream transcription factor 2, c-fos interacting	X90824	1.35
KIAA0268 protein	AK025122	1.35
f-box and leucine-rich repeat protein 4	AF176699	1.35
tyrosine 3-monooxygenase/tryptophan 5-monooxygenase activation protein, epsilon polypeptide	U43430	1.35
ribosomal protein L31	NM_000993	1.35
Consensus includes gb:AA721025 /FEA=EST /DB_XREF=gi:2737160 /DB_XREF=est:nz89g05.s1 /CLONE=IMAGE:1269464 /UG=Hs.293253 ESTs	AA721025	1.35
translocase of inner mitochondrial membrane 44 homolog (yeast)	AF026030	1.35
v-crk sarcoma virus CT10 oncogene homolog (avian)	NM_016823	1.35
defensin, beta 2	NM_004942	1.35
translocase of inner mitochondrial membrane 22 homolog (yeast)	NM_013337	1.35
KIAA0240 protein	AI801951	1.35
putative G-protein coupled receptor	NM_016334	1.35
IGF-II mRNA-binding protein 3	AL023775	1.35
SELENOPHOSPHATE SYNTHETASE ; Human selenium donor protein	AV682679	1.35
protein kinase C, delta	NM_006254	1.35
uncharacterized hypothalamus protein HARP11	AK022248	1.35
karyopherin (importin) beta 3	AU148466	1.35
thyroid hormone receptor-associated protein, 95-kD subunit	AA314406	1.35
hypothetical protein FLJ13612	BC002449	1.35
dynein, cytoplasmic, light polypeptide	NM_003746	1.35
BAI1-associated protein 3	AI799802	1.35
small nuclear RNA activating complex, polypeptide 3, 50kD	U71300	1.35
Wolf-Hirschhorn syndrome candidate 2	NM_005663	1.35
HSC6.1.1 protein	AB016900	1.35
KIAA0355 gene product	NM_014686	1.35
YY1 transcription factor	Z14077	1.35
ras homolog gene family, member C	NM_005167	1.35
DKFZP564A043 protein	AL036344	1.35
casein kinase 1, alpha 1	AW268585	1.35
hypothetical protein	AF151022	1.35
general transcription factor II, i	AC004883	1.35
uncharacterized hematopoietic stem/progenitor cells protein MDS028	NM_018463	1.35
splicing factor 1	NM_004630	1.35
GATA binding protein 3	AI796169	1.35
Consensus includes gb:AW575379 /FEA=EST /DB_XREF=gi:7246918 /DB_XREF=est:UI-HF-BL0-adc-e-05-0-UI.s1 /CLONE=IMAGE:3060969 /UG=Hs.6551 ATPase, H+ transporting, lysosom	AW575379	1.35
anaphase-promoting complex subunit 10	NM_014885	1.35
hypothetical protein FLJ21877	NM_022065	1.35
Consensus includes gb:AW128846 /FEA=EST /DB_XREF=gi:6116764 /DB_XREF=est:xd46f06.x1 /CLONE=IMAGE:2596835 /UG=Hs.244737 ESTs, Weakly similar to neuronal thread protein /	AW128846	1.35
HSPC159 protein	NM_014181	1.35
RNA binding motif, single stranded interacting protein 1	NM_016839	1.35
hypothetical protein R33729_1	Z78330	1.35
potassium channel modulatory factor	NM_020122	1.35

hypothetical protein	NM_018988	1.35
6 protein-coupled receptor 38	NM_001507	1.35
hypothetical protein FLJ10652	NM_018169	1.35
heterogeneous nuclear ribonucleoprotein A3	BG505670	1.35
intersectin 2	AF182198	1.35
tetratricopeptide repeat domain 3	D83077	1.35
nuclear pore complex interacting protein	AF229069	1.35
tumor necrosis factor, alpha-induced protein 2	AI862445	1.35
ELK3, ETS-domain protein (SRF accessory protein 2)	AW575374	1.35
hypothetical protein FLJ10154	NM_018011	1.35
chromosome 5 open reading frame 6	NM_016605	1.35
UDP-N-acteylglucosamine pyrophosphorylase 1	S73498	1.35
ATP citrate lyase	AI971281	1.35
nuclear cap binding protein subunit 2, 20kD	BC001255	1.35
hypothetical protein FLJ23459	NM_024775	1.35
FUS interacting protein (serine-arginine rich) 2	NM_021993	1.35
t-complex-associated-testis-expressed 1-like 1	NM_006519	1.35
SCN Circadian Oscillatory Protein (SCOP)	AB011178	1.35
DKFZP58681621 protein	NM_015533	1.35
cytochrome b-561	BC002976	1.35
KIAA0648 protein	AW991219	1.34
MYB binding protein (P160) 1a	NM_014520	1.34
zinc finger RNA binding protein	BC000376	1.34
abl-interactor 12 (SH3-containing protein)	AF260261	1.34
hypothetical protein FLJ10709	NM_018188	1.34
KIAA0592 protein	BC006456	1.34
vascular endothelial growth factor	M27281	1.34
hypothetical protein FLJ22222	W19668	1.34
calcium channel, voltage-dependent, gamma subunit 2	AL022313	1.34
transcriptional intermediary factor 1	AL538264	1.34
KIAA0810 protein	BE615699	1.34
Consensus includes gb:AA928506 /FEA=EST /DB_XREF=gi:3076797 /DB_XREF=est:om17g03.s1 /CLONE=IMAGE:1541332 /UG=Hs.10762 ESTs	AA928506	1.34
galanin receptor 2	AF080586	1.34
ribosomal protein L5	AK027146	1.34
Consensus includes gb:BE466525 /FEA=EST /DB_XREF=gi:9512223 /DB_XREF=est:hx94b10.x1 /CLONE=IMAGE:3195451 /UG=Hs.234773 Homo sapiens cDNA: FLJ22281 fis, clone HRC03:	BE466525	1.34
splicing factor, arginine/serine-rich 4	NM_005626	1.34
ems1 sequence (mammary tumor and squamous cell carcinoma-associated (p80/85 src substrate)	BG475299	1.34
Consensus includes gb:J04742 /DEF=Human autonomous replicating sequence H1 (ARSH1) /FEA=CDS /DB_XREF=gi:179028 /UG=Hs.247945 Human autonomous replicating sequence H1	J04742	1.34
chromosome 11 open reading frame 24	NM_022338	1.34
potassium voltage-gated channel, shaker-related subfamily, beta member 1	U16953	1.34
hypothetical protein FLJ20727	BE966019	1.34
agrin	AK021586	1.34
etoposide-induced mRNA	NM_004879	1.34
Consensus includes gb:BE869922 /FEA=EST /DB_XREF=gi:10318698 /DB_XREF=est:601446568F1 /CLONE=IMAGE:3850432 /UG=Hs.181307 H3 histone, family 3A	BE869922	1.34
KIAA0676 protein	AK026096	1.34
KIAA0210 gene product	AI300084	1.34
peroxisomal D3,D2-enoyl-CoA isomerase	NM_006117	1.34
Consensus includes gb:AL049250.1 /DEF=Homo sapiens mRNA; cDNA DKFZp564D113 (from clone DKFZp564D113). /FEA=mRNA /DB_XREF=gi:4499989 /UG=Hs.194637 BANP homolog, S	AL049250	1.34
Sec23-interacting protein p125	AK001135	1.34
cytochrome P450, subfamily I (dioxin-inducible), polypeptide 1 (glaucoma 3, primary infantile)	N21019	1.34
lectin, galactoside-binding, soluble, 3 (galectin 3)	BC001120	1.34
citron (rho-interacting, serine/threonine kinase 21)	AI861788	1.34
rab6 GTPase activating protein (GAP and centrosome-associated)	NM_012197	1.34
LCAT-like lysophospholipase	AL110209	1.34
HCR (a-helix coiled-coil rod homologue)	AB029343	1.34
v-myc myelocytomatosis viral oncogene homolog (avian)	NM_002467	1.34
Consensus includes gb:AW409974 /FEA=EST /DB_XREF=gi:6935515 /DB_XREF=est:fh03d12.y1 /CLONE=IMAGE:2961094 /UG=Hs.332750 Homo sapiens cDNA FLJ10100 fis, clone HEMBA	AW409974	1.34
NS1-binding protein	AF205218	1.34
ribosomal protein L37a	NM_000998	1.34
KIAA1097 protein	AB029020	1.34
RNA binding motif protein 15	NM_022768	1.34
hypothetical protein MGC3077	AK021779	1.34
VRK3 for vaccinia related kinase 3	NM_016440	1.34

progesterone receptor membrane component 1	NM_006667	1.34
uncharacterized hypothalamus protein HT007	AF246240	1.34
plexin B2	BC004542	1.34
uncharacterized bone marrow protein BM039	AK023669	1.34
DNA segment on chromosome 12 (unique) 2489 expressed sequence	NM_007360	1.34
Consensus includes gb:AI742553 /FEA=EST /DB_XREF=gi:5110841 /DB_XREF=est:wg55d02.x1 /CLONE=IMAGE:2368995 /UG=Hs.184592 protein kinase, lysine deficient 1 /FL=gb:NM_01	AI742553	1.34
Consensus includes gb:W84525 /FEA=EST /DB_XREF=gi:1395637 /DB_XREF=est:zd90g09.s1 /CLONE=IMAGE:356800 /UG=Hs.7949 DKFZP586B2420 protein	W84525	1.34
neutral sphingomyelinase (N-SMase) activation associated factor	NM_003580	1.34
CGI-92 protein	NM_016035	1.34
2',5'-oligoadenylate synthetase 2 (69-71 kD)	NM_016817	1.34
pazillin	D86862	1.34
hypothetical protein FLJ22195	NM_022758	1.34
KIAA0586 gene product	AB011158	1.34
ortholog of mouse integral membrane glycoprotein LIG-1	AB050468	1.34
NME7	NM_013330	1.34
G protein-coupled receptor 43	NM_005306	1.34
protein tyrosine phosphatase, receptor type, f polypeptide (PTPRF), interacting protein (liprin), alpha 1	AA195259	1.34
Consensus includes gb:BF057059 /FEA=EST /DB_XREF=gi:10810955 /DB_XREF=est:7k15c07.x1 /CLONE=IMAGE:3444468 /UG=Hs.12421 hypothetical protein	BF057059	1.34
Consensus includes gb:BF056534 /FEA=EST /DB_XREF=gi:10810430 /DB_XREF=est:7k06e09.x1 /CLONE=IMAGE:3443441 /UG=Hs.123469 ESTs, Weakly similar to AF208855 1 BM-013 H	BF056534	1.34
homolog of rat nadrin	NM_018054	1.33
lamin B1	NM_005573	1.33
small GTP-binding protein	NM_030981	1.33
amyloid beta precursor protein binding protein 1, 59kD	NM_003905	1.33
general transcription factor IIIA	NM_002097	1.33
homolog of mouse quaking QKI (KH domain RNA binding protein)	AF142418	1.33
calcium channel, voltage-dependent, gamma subunit 2	AL022313	1.33
platelet derived growth factor C	NM_016205	1.33
phosphoribosyl pyrophosphate synthetase-associated protein 2	NM_002767	1.33
stromal antigen 1	AI126490	1.33
KIAA0737 gene product	BE783632	1.33
lactotransferrin	NM_002343	1.33
gp25L2 protein	BC001123	1.33
KIAA0747 protein	BC004998	1.33
RAN binding protein 2	NM_006267	1.33
lactate dehydrogenase B	BE042354	1.33
transcription factor 2, hepatic; LF-B3; variant hepatic nuclear factor	NM_000458	1.33
Fanconi anemia, complementation group G	NM_004629	1.33
ribosomal protein L9	NM_000661	1.33
clones 23667 and 23775 zinc finger protein	BC000330	1.33
glycosylphosphatidylinositol specific phospholipase D1	AL031230	1.33
destrin (actin depolymerizing factor)	NM_006870	1.33
cysteine and glycine-rich protein 1	NM_004078	1.33
protein tyrosine kinase 2 beta	U43522	1.33
protease, cysteine, 1 (legumain)	D55696	1.33
methyl-CpG binding domain protein 1	NM_015846	1.33
mitogen-activated protein kinase kinase 2	AI762811	1.33
nuclear pore complex interacting protein	AC002045	1.33
adenylate kinase 2	AU154985	1.33
Consensus includes gb:AK022459.1 /DEF=Homo sapiens cDNA FLJ12397 fis, clone MAMMA1002769, weakly similar to Homo sapiens cell cycle progression restoration 8 protein (CPR8) mRN	AK022459	1.33
NAD kinase	BE674658	1.33
flightless I homolog (Drosophila)	BG421186	1.33
SIPL protein	NM_018269	1.33
myosin, light polypeptide 4, alkali; atrial, embryonic	X58851	1.33
TAR DNA binding protein	NM_007375	1.33
DKFZP434D193 protein	AL080129	1.33
serologically defined colon cancer antigen 3	NM_006643	1.33
branched chain keto acid dehydrogenase E1, beta polypeptide (maple syrup urine disease)	M55575	1.33
hypothetical protein FLJ13868	NM_022744	1.33
SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily a, member 2	NM_003070	1.33
MCM6 minichromosome maintenance deficient 6 (MIS5 homolog, S. pombe) (S. cerevisiae)	NM_005915	1.33
TIA1 cytotoxic granule-associated RNA binding protein	AL046419	1.33
calnexin	M94859	1.33
hypothetical protein dJ465N24.2.1	AF247168	1.33

calcium/calmodulin-dependent protein kinase kinase 2, beta	AK024748	1.33
arylsulfatase E (chondrodysplasia punctata 1)	NM_000047	1.33
MYC-associated zinc finger protein (purine-binding transcription factor)	AI471665	1.33
origin recognition complex, subunit 3-like (yeast)	AF125507	1.33
Consensus includes gb:AL049314.1 /DEF=Homo sapiens mRNA; cDNA DKFZp564C156 (from clone DKFZp564C156). /FEA=mRNA /DB_XREF=gi:4500088 /UG=Hs.278592 Homo sapiens mR	AL049314	1.33
RAP1, GTPase activating protein 1	NM_002885	1.33
protein tyrosine phosphatase, receptor type, N	NM_002846	1.33
zinc finger protein 36, C3H type-like 2	U07802	1.33
X-box binding protein 1	NM_005080	1.33
hypothetical protein FLJ10734	NM_018197	1.33
Consensus includes gb:AJ295618 /DEF=Homo sapiens FTSH gene for putative ATPases, exons 1 and 2 and join CDS /FEA=mRNA /DB_XREF=gi:9506352 /UG=Hs.206521 YME1 (S.cerevisia	AJ295618	1.33
ubiquitin C-terminal hydrolase UCH37	NM_015984	1.33
gb:NM_002913.1 /DEF=Homo sapiens replication factor C (activator 1) 1 (145kD) (RFC1), mRNA. /FEA=mRNA /GEN=RFC1 /PROD=replication factor C (activator 1) 1 (145kD) /DB_XREF=	NM_002913	1.33
copine VI (neuronal)	AB009288	1.33
Williams-Beuren syndrome chromosome region 1	NM_022170	1.33
polymerase (DNA directed), mu	AK023002	1.33
kinesin-like 2	BC000712	1.33
nudix (nucleoside diphosphate linked moiety X)-type motif 2	NM_001161	1.33
KIAA1624 protein	AK024651	1.33
activin A receptor, type IIB	NM_001106	1.33
mutL homolog 1, colon cancer, nonpolyposis type 2 (E. coli)	NM_000249	1.33
tumor protein, translationally-controlled 1	AI888178	1.33
prp28, U5 snRNP 100 kd protein	AF026402	1.33
peroxisomal biogenesis factor 11B	NM_003846	1.33
KIAA0701 protein	AL136821	1.33
COX15 homolog, cytochrome c oxidase assembly protein (yeast)	NM_004376	1.32
chromosome 8 open reading frame 2	T90773	1.32
cortistatin	AL578583	1.32
CDC28 protein kinase 1	NM_001826	1.32
regulator of nonsense transcripts 1	U59323	1.32
KIAA1449 protein	N64681	1.32
opioid growth factor receptor	AF172452	1.32
MYLE protein	NM_014015	1.32
TIA1 cytotoxic granule-associated RNA binding protein-like 1	NM_003252	1.32
DKFZP564C186 protein	NM_015658	1.32
MAGE-E1 protein	NM_030801	1.32
protocadherin LKC	NM_017675	1.32
forkhead box H1	NM_003923	1.32
Consensus includes gb:AI472139 /FEA=EST /DB_XREF=gi:4334229 /DB_XREF=est:tj86a10.x1 /CLONE=IMAGE:2148378 /UG=Hs.9683 DnaJ (Hsp40) homolog, subfamily C, member 3	AI472139	1.32
hypothetical protein FLJ22427	BF129339	1.32
S-adenosylhomocysteine hydrolase-like 1	AA479488	1.32
Sec23-interacting protein p125	AK021846	1.32
nuclear pore complex interacting protein	AA308853	1.32
CGI-82 protein	AF167438	1.32
baculoviral IAP repeat-containing 7 (livin)	NM_022161	1.32
hypothetical protein	NM_015702	1.32
protein tyrosine phosphatase, non-receptor type 2	AI828880	1.32
malate dehydrogenase 2, NAD (mitochondrial)	AL520774	1.32
KIAA0014 gene product	NM_014665	1.32
pyruvate dehydrogenase (lipoamide) beta	AL117618	1.32
mitochondrial ribosomal protein S18A	AB049952	1.32
protein phosphatase 3 (formerly 2B), catalytic subunit, alpha isoform (calcineurin A alpha)	AL353950	1.32
6-pyruvoyltetrahydropterin synthase	M97655	1.32
olfactory receptor, family 2, subfamily B, member 6	AL133267	1.32
gastrin-releasing peptide receptor	NM_005314	1.32
RAS guanyl releasing protein 2 (calcium and DAG-regulated)	NM_005825	1.32
hypothetical protein FLJ11149	BF340123	1.32
Consensus includes gb:BE856549 /FEA=EST /DB_XREF=gi:10369681 /DB_XREF=est:7164b02.x1 /CLONE=IMAGE:3299403 /UG=Hs.44131 KIAA0974 protein	BE856549	1.32
SET translocation (myeloid leukemia-associated)	AV702810	1.32
phosphatase and tensin homolog (mutated in multiple advanced cancers 1)	U96180	1.32
transmembrane 9 superfamily member 1	U94831	1.32
TNFRSF1A-associated via death domain	L41690	1.32
KIAA0467 protein	NM_024547	1.32

neurochondrin	AB011179	1.32
phosphotidylinositol transfer protein, beta	AL031591	1.32
hypothetical protein FLJ22028	AI614728	1.32
ATP synthase, H+ transporting, mitochondrial F0 complex, subunit c (subunit 9), isoform 2	X69909	1.32
Consensus includes gb:L10374.1 /DEF=Human (clone CTG-A4) mRNA sequence. /FEA=mRNA /DB_XREF=gi:307288 /UG=Hs.281587 Human (clone CTG-A4) mRNA sequence	L10374	1.32
quinolinate phosphoribosyltransferase (nicotinate-nucleotide pyrophosphorylase (carboxylating))	NM_014298	1.32
FK506 binding protein like	NM_022110	1.32
hsp70-interacting protein	NM_012267	1.32
f-box and leucine-rich repeat protein 11	BE675843	1.32
homolog of mouse quaking QKI (KH domain RNA binding protein)	AA149639	1.32
hypothetical protein FLJ10914	NM_018270	1.32
peroxisomal biogenesis factor 14	NM_004565	1.32
hypothetical protein FLJ13409	NM_024617	1.32
hypothetical protein FLJ22622	NM_025151	1.32
arfaptin 1	AL044651	1.32
cornichon-like	NM_005776	1.32
nicastrin	AF240468	1.32
glucose-6-phosphatase, transport (glucose-6-phosphate) protein 1	NM_001467	1.32
TRAM-like protein	AI986461	1.32
KIAA0217 protein	AW628686	1.32
integrin beta 3 binding protein (beta3-endonexin)	NM_014288	1.32
FAST kinase	NM_006712	1.32
hypothetical protein FLJ22578	NM_024864	1.32
G protein pathway suppressor 2	AL157493	1.32
U2 small nuclear ribonucleoprotein auxiliary factor (65kD)	NM_007279	1.32
phosphoribosylglycinamide formyltransferase, phosphoribosylglycinamide synthetase, phosphoribosylaminoimidazole synthetase	AF008655	1.32
hypothetical protein FLJ10439	NM_018093	1.32
KIAA0725 protein	AB018268	1.32
Consensus includes gb:AL078596 /DEF=Human DNA sequence from clone RP3-429G5 on chromosome 6q21-22.1. Contains the NR2E1 gene for nuclear receptor 2E1 (tailless, TLL, TLX, XTL	AL078596	1.32
KIAA1035 protein	NM_015239	1.32
SH3-domain GRB2-like 1	NM_003025	1.32
microsomal glutathione S-transferase 2	NM_002413	1.32
nuclear receptor subfamily 5, group A, member 2	AF146343	1.32
C2H2 (Kruppel-type) zinc finger protein	AF159567	1.32
cofactor required for Sp1 transcriptional activation, subunit 2 (150kD)	AI971089	1.32
ATPase, H+ transporting, lysosomal (vacuolar proton pump), alpha polypeptide, 70kD, isoform 1	NM_001690	1.32
FLN29 gene product	AB007447	1.32
B-cell CLL/lymphoma 7A	NM_020993	1.32
oxysterol binding protein-like 3	AI202969	1.31
hypothetical protein FLJ21749	NM_025124	1.31
SPTF-associated factor 65 gamma	AF197954	1.31
transforming growth factor, beta receptor II (70-80kD)	D50683	1.31
diptheria toxin resistance protein required for diphthamide biosynthesis (Saccharomyces)-like 2	NM_001384	1.31
phosphatidylinositol transfer protein, membrane-associated	NM_004910	1.31
glycerol kinase	NM_000167	1.31
CD63 antigen (melanoma 1 antigen)	NM_001780	1.31
hypothetical protein FLJ12428	NM_022783	1.31
hypothetical protein MGC4504	NM_024111	1.31
Ac-like transposable element	NM_004729	1.31
choline kinase	AI991328	1.31
ribosomal protein L13	AA961748	1.31
rTS beta protein	NM_017512	1.31
pleiomorphic adenoma gene-like 2	NM_002657	1.31
Treacher Collins-Franceschetti syndrome 1	NM_000356	1.31
RAE1 RNA export 1 homolog (S. pombe)	NM_003610	1.31
Consensus includes gb:BG178274 /FEA=EST /DB_XREF=gi:12684977 /DB_XREF=est:602330318F1 /CLONE=IMAGE:4431676 /UG=Hs.278895 postmeiotic segregation increased 2-like 5	BG178274	1.31
poliovirus receptor-related 1 (herpesvirus entry mediator C; nectin)	NM_002855	1.31
uncharacterized hematopoietic stem/progenitor cells protein MDS033	NM_018468	1.31
CGI-74 protein	NM_016019	1.31
mitogen-activated protein kinase kinase kinase kinase 5	NM_006575	1.31
splicing factor 3b, subunit 1, 155kD	AI739389	1.31
Consensus includes gb:AI375965 /FEA=EST /DB_XREF=gi:4175955 /DB_XREF=est:tc24h12.s1 /CLONE=IMAGE:2065607 /UG=Hs.131067 ESTs, Moderately similar to IBA2_HUMAN ITBA2 P	AI375965	1.31
colony stimulating factor 1 (macrophage)	NM_000757	1.31

Fig

endocytic receptor (macrophage mannose receptor family)	AB014609	1.31
ATPase, Ca++ transporting, ubiquitous	NM_005173	1.31
mitochondrial ribosomal protein 63	BF303597	1.31
ubiquitin specific protease 12	AF022790	1.31
Consensus includes gb:AL109722.1 /DEF=Homo sapiens mRNA full length insert cDNA clone EUROIMAGE 31619. /FEA=mRNA /DB_XREF=gi:5689814 /UG=Hs.170079 Homo sapiens mRNA	AL109722	1.31
hypothetical protein FLJ13491	AJ309636	1.31
CGI-01 protein	AL049669	1.31
BUB1 budding uninhibited by benzimidazoles 1 homolog (yeast)	AF043294	1.31
transposon-derived Buster3 transposase-like	NM_022090	1.31
leptin receptor overlapping transcript-like 1	NM_015344	1.31
hypothetical protein FLJ22127	AI028241	1.31
ribosomal protein L38	NM_000999	1.31
ribosomal protein S6	NM_001010	1.31
Down syndrome critical region gene 3	NM_006052	1.31
SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily d, member 2	AF113019	1.31
stromal antigen 2	BC001765	1.31
inositol polyphosphate-4-phosphatase, type I, 107kD	NM_001566	1.31
calcium channel, voltage-dependent, beta 1 subunit	M92301	1.31
brain and reproductive organ-expressed (TNFRSF1A modulator)	NM_004899	1.31
aldehyde dehydrogenase 5 family, member A1 (succinate-semialdehyde dehydrogenase)	NM_001080	1.31
hypothetical protein RG083M05.2	AK026142	1.31
hypothetical protein FLJ10719	BG403615	1.31
hypothetical protein	BC001393	1.31
Tax interaction protein 1	AF234997	1.31
seven transmembrane domain orphan receptor	NM_016372	1.31
core-binding factor, beta subunit	NM_001755	1.31
ceroid-lipofuscinosis, neuronal 2, late infantile (Jansky-Bielschowsky disease)	NM_000391	1.31
FAST kinase	AK023141	1.31
hypothetical protein	AA463853	1.31
similar to hypothetical protein MNCb-2386	BG230758	1.31
U6 snRNA-associated Sm-like protein LSm8	NM_016200	1.31
Bardet-Biedl syndrome 4	AI813772	1.31
chromosome 21 open reading frame 45	NM_018944	1.31
phosphotidylinositol transfer protein	H15647	1.31
SELENOPHOSPHATE SYNTHETASE ; Human selenium donor protein	AI885670	1.31
2'-5'-oligoadenylate synthetase 2 (69-71 kD)	NM_002535	1.31
polymerase I and transcript release factor	AF312393	1.31
heterogeneous nuclear ribonucleoprotein A0	NM_006805	1.31
flotillin 2	BC003683	1.31
Consensus includes gb:AL522296 /FEA=EST /DB_XREF=gi:12785789 /DB_XREF=est:AL522296 /CLONE=CS0DB008YP08 (5 prime) /UG=Hs.17481 Homo sapiens clone 24606 mRNA sequ	AL522296	1.31
Consensus includes gb:AW081820 /FEA=EST /DB_XREF=gi:6036972 /DB_XREF=est:xb55d12.x1 /CLONE=IMAGE:2580215 /UG=Hs.287317 ESTs, Weakly similar to A59252 myosin heavy cl	AW081820	1.31
IM5 protein	BC000464	1.31
hypothetical protein FLJ20203	NM_017710	1.31
GRB2-associated binding protein 1	NM_002039	1.31
hypothetical protein FLJ10737	AL109978	1.31
calpain 1, (mu/I) large subunit	NM_005186	1.31
Consensus includes gb:M22005 /DEF=Human interleukin 2 gene, clone pATtacIL-2C2TT, complete cds, clone pATtacIL-2C2TT /FEA=CDS /DB_XREF=gi:186300 /UG=Hs.247956 Human interl	M22005	1.31
Cluster Incl. AA522670:ni39a05.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-979184 /clone_end=3 /gb=AA522670 /gi=2263382 /ug=Hs.7247 /len=684	AA522670	1.31
KIAA1232 protein	AB033058	1.31
glutaminase	NM_014905	1.31
hypothetical protein FLJ22794	NM_022074	1.31
transporter 1, ATP-binding cassette, sub-family B (MDR/TAP)	NM_000593	1.30
KIAA0176 protein	AA523921	1.30
high-mobility group 20A	NM_018200	1.30
DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 28	AC004531	1.30
hypothetical protein FLJ10283	NM_018046	1.30
KIAA0483 protein	AB007952	1.30
hypothetical protein FLJ20477	NM_017837	1.30
cathelicidin antimicrobial peptide	U19970	1.30
Bruton agammaglobulinemia tyrosine kinase	NM_000061	1.30
KIAA0447 gene product	BE885244	1.30
poly(A) binding protein, cytoplasmic 4 (inducible form)	NM_003819	1.30
dopamine receptor D2	M30625	1.30

RNA binding protein S1, serine-rich domain	BC001659	1.30
copine III	AA541758	1.30
N-acetylglucosamine-phosphate mutase	BC001258	1.30
chromosome 9 open reading frame 12	NM_022755	1.30
hypothetical protein FLJ10767	NM_018209	1.30
hypothetical protein FLJ22624	AW043622	1.30
hypothetical protein MGC5466	AI681536	1.30
cell division cycle 42 (GTP binding protein, 25kD)	R37664	1.30
activated p21cdc42Hs kinase	NM_005781	1.30
Consensus includes gb:AI652058 /FEA=EST /DB_XREF=gi:4736037 /DB_XREF=est:wb27d11.x1 /CLONE=IMAGE:2306901 /UG=Hs.129061 Human Chromosome 16 BAC clone CIT987SK-A-1	AI652058	1.30
valyl-tRNA synthetase 2	BE790854	1.30
hypothetical protein FLJ10698	AF062733	1.30
golgi phosphoprotein 5	AK001574	1.30
crystallin, zeta (quinone reductase)-like 1	NM_005111	1.30
zinc finger protein 132 (clone pHZ-12)	NM_003433	1.30
Down syndrome critical region gene 5	AB035745	1.30
topoisomerase (DNA) II alpha (170kD)	AU159942	1.30
pumilio (Drosophila) homolog 2	BE778078	1.30
small nuclear ribonucleoprotein D3 polypeptide (18kD)	NM_004175	1.30
Consensus includes gb:AK025086.1 /DEF=Homo sapiens cDNA: FLJ21433 fis, clone COL04232. /FEA=mRNA /DB_XREF=gi:10437530 /UG=Hs.306761 Homo sapiens cDNA: FLJ21433 fis, cl	AK025086	1.30
KIAA0176 protein	D79998	1.30
mitogen-activated protein kinase 1	NM_002745	1.30
chromosome 11 open reading frame 23	NM_018312	1.30
Ts translation elongation factor, mitochondrial	AF110399	1.30
hypothetical protein FLJ21940	AW975818	1.30
hypothetical protein FLJ22028	AI962943	1.30
N51-associated protein 1	AF037448	1.30
protocadherin 9	AI524125	1.30
copine I	NM_003915	1.30
tumor rejection antigen (gp96) 1	NM_003299	1.30
protein tyrosine phosphatase, receptor type, C-associated protein	NM_005608	1.30
proteasome (prosome, macropain) subunit, beta type, 1	W86293	1.30
Consensus includes gb:AV661533 /FEA=EST /DB_XREF=gi:9882547 /DB_XREF=est:AV661533 /CLONE=GLCGTD12 /UG=Hs.293637 ESTs	AV661533	1.30
signal-induced proliferation-associated gene 1	NM_006747	1.30
cytochrome c oxidase subunit Va	NM_004255	1.30
eukaryotic translation initiation factor 4 gamma, 2	NM_001418	1.30
protein phosphatase 2 (formerly 2A), regulatory subunit A (PR 65), beta isoform	T79584	1.30
O-linked N-acetylglucosamine (GlcNAc) transferase (UDP-N-acetylglucosamine:polypeptide-N-acetylglucosaminyl transferase)	AF070560	1.30
KIAA0008 gene product	NM_014750	1.30
NAD(P)H dehydrogenase, quinone 2	NM_000904	1.30
actin cross-linking factor	AK023406	1.30
Ras-GTPase activating protein SH3 domain-binding protein 2	AB014560	1.30
dynactin 2 (p50)	NM_006400	1.30
tripartite motif-containing 26	NM_003449	1.30
Sec23-interacting protein p125	AI332962	1.30
delta-tubulin	BC000258	1.30
Rag C protein	NM_022157	1.30
death effector domain-containing	AF064605	1.30
acetylserotonin O-methyltransferase-like	AA669799	1.30
phosphofructokinase, muscle	U24183	1.30
flotillin 1	AF085357	1.30
leucine aminopeptidase	NM_015907	1.30
KIAA0690 protein	AK021460	1.30
KIAA1089 protein	AB029012	1.30
hypothetical protein	BC002568	1.30
super conserved receptor expressed in brain 3	NM_018969	1.30
Notch homolog 2 (Drosophila)	AA291203	1.30
decorin	NM_001920	1.30
KIAA0999 protein	AA044154	1.30
cholinergic receptor, nicotinic, alpha polypeptide 7	U62436	1.30
6 protein-binding protein CRF6	NM_012341	1.30
CGI-05 protein	NM_016408	1.30
Consensus includes gb:AC004490 /DEF=Homo sapiens chromosome 19, cosmid R29381 /FEA=CDS /DB_XREF=gi:2995576 /UG=Hs.156015 Homo sapiens chromosome 19, cosmid R29381	AC004490	1.30

endocrine regulator	NM_014345	1.30
superoxide dismutase 2, mitochondrial	W46388	1.30
zinc finger protein 354A	NM_005649	1.30
KIAA0561 protein	AB011133	1.29
Consensus includes gb:Y15724 /DEF=Homo sapiens SERCA3 gene, exons 1-7 (and joined CDS) /FEA=mRNA /DB_XREF=gi:3021395 /UG=Hs.5541 ATPase, Ca++ transporting, ubiquitous	Y15724	1.29
protein phosphatase 4, regulatory subunit 1	NM_005134	1.29
KIAA0871 protein	AF112221	1.29
thioredoxin-like 2	AL138831	1.29
proteasome (prosome, macropain) subunit, alpha type, 2	AL523904	1.29
growth arrest and DNA-damage-inducible, beta	NM_015675	1.29
MCT-1 protein	NM_014060	1.29
methyl-CpG binding domain protein 4	AL556619	1.29
phosphoinositide-3-kinase, catalytic, delta polypeptide	U86453	1.29
hypothetical protein FLJ22637	NM_025165	1.29
parathymosin	M24398	1.29
vimentin	AI922599	1.29
calreticulin	AI348935	1.29
ATPase, H+ transporting, lysosomal (vacuolar proton pump), beta polypeptide, 56/58kD, isoform 1 (Renal tubular acidosis with deafness)	NM_001692	1.29
Bicaudal D homolog 1 (Drosophila)	U90030	1.29
gene predicted from cDNA with a complete coding sequence	NM_014628	1.29
inositol (myo)-1(or 4)-monophosphatase 1	NM_005536	1.29
Consensus includes gb:AL161956.1 /DEF=Homo sapiens mRNA; cDNA DKFZp761A17121 (from clone DKFZp761A17121). /FEA=mRNA /DB_XREF=gi:7328008 /UG=Hs.306492 Homo sapie	AL161956	1.29
ribosomal protein L13a	BC000514	1.29
SA (rat hypertension-associated) homolog	AC004381	1.29
splicing factor (CC1.3)	NM_004902	1.29
BCR downstream signaling 1	NM_012108	1.29
lymphocyte adaptor protein	NM_005475	1.29
karyopherin (importin) beta 1	L38951	1.29
homolog of yeast ribosome biogenesis regulatory protein RRS1	BC001811	1.29
hypothetical protein FLJ20186	NM_017702	1.29
hypothetical protein MGC4126	AI357539	1.29
apolipoprotein M	NM_019101	1.29
inhibitor of DNA binding 2, dominant negative helix-loop-helix protein	D13891	1.29
diazepam binding inhibitor (GABA receptor modulator, acyl-Coenzyme A binding protein)	BC006466	1.29
lysosomal-associated membrane protein 1	J03263	1.29
KIAA1332 protein	AA836114	1.29
RNA binding motif, single stranded interacting protein 1	NM_016837	1.29
hypothetical protein DKFZp564O1664	NM_030800	1.29
Consensus includes gb:AL137398.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434K052 (from clone DKFZp434K052). /FEA=mRNA /DB_XREF=gi:6807944 /UG=Hs.169639 Homo sapiens mR	AL137398	1.29
plakophilin 4	BG292559	1.29
DnaJ (Hsp40) homolog, subfamily B, member 9	NM_012328	1.29
hypothetical protein FLJ21736	NM_024922	1.29
BUB3 budding uninhibited by benzimidazoles 3 homolog (yeast)	NM_004725	1.29
Consensus includes gb:AW293356 /FEA=EST /DB_XREF=gi:6699992 /DB_XREF=est:UI-H-BI2-ahl-c-11-0-UI.s1 /CLONE=IMAGE:2727165 /UG=Hs.58220 Homo sapiens cDNA: FLJ23005 fis,	AW293356	1.29
Consensus includes gb:AF258545 /DEF=Homo sapiens chormosome 17, HC90, HC71AC, HC6 and HC56 genes, complete sequence /FEA=CDS /DB_XREF=gi:10945429 /UG=Hs.19614 gemir	AF258545	1.29
Wilms' tumour 1-associating protein	NM_004906	1.29
guanine nucleotide binding protein (G protein), alpha z polypeptide	NM_002073	1.29
BRCA1 associated protein	AL042733	1.29
ferredoxin reductase	NM_004110	1.29
MADS box transcription enhancer factor 2, polypeptide C (myocyte enhancer factor 2C)	N22468	1.29
KIAA0682 gene product	NM_016196	1.29
glyoxalase I	NM_006708	1.29
RAD51 homolog (RecA homolog, E. coli) (S. cerevisiae)	NM_002875	1.29
chromosome 6 open reading frame 34	NM_014320	1.29
NCK adaptor protein 1	NM_006153	1.29
clathrin, light polypeptide (Lcb)	BC006332	1.29
glycoprotein V (platelet)	NM_004488	1.29
hypothetical protein FLJ20154	AK000161	1.29
Consensus includes gb:AW449390 /FEA=EST /DB_XREF=gi:6990166 /DB_XREF=est:UI-H-BI3-aki-g-10-0-UI.s1 /CLONE=IMAGE:2734650 /UG=Hs.257150 ESTs, Moderately similar to SUR1,	AW449390	1.29
DKFZP564O092 protein	AI269117	1.29
KIAA0543 protein	AB011115	1.29
ras-like protein	NM_012249	1.29
GLUT4 enhancer factor	NM_020062	1.29

angiopoietin-3	NM_015985	1.29
6-phosphofructo-2-kinase/fructose-2,6-biphosphatase 3	NM_004566	1.29
DnaJ (Hsp40) homolog, subfamily B, member 1	NM_006145	1.29
hypothetical protein FLJ12847	NM_024596	1.29
phospholipase C, epsilon 2	AL117515	1.29
hypothetical protein MGC10882	BC004952	1.29
fibroblast growth factor 5	AB016517	1.29
translocating chain-associating membrane protein	AF130055	1.29
cleavage and polyadenylation specific factor 6, 68kD subunit	AU149367	1.29
peroxisomal biogenesis factor 3	AB035307	1.29
translocase of inner mitochondrial membrane 23 homolog (yeast)	NM_006327	1.29
CASP8 and FADD-like apoptosis regulator	AF015451	1.29
putative methyltransferase	AW069290	1.29
hypothetical protein FLJ23018	NM_024810	1.29
H1 histone family, member 0	BC000145	1.29
spastic paraplegia 7, paraplegin (pure and complicated autosomal recessive)	NM_003119	1.29
protein tyrosine phosphatase type IVA, member 2	U48297	1.29
Rho GTPase activating protein 1	BG468434	1.29
RNA binding motif, single stranded interacting protein 1	D28482	1.29
gb:NM_015870.1 /DEF=Homo sapiens endogenous retrovirus H D1 leader regionintegrase-derived ORF1, ORF2, and putative envelope protein (HSU88895), mRNA. /FEA=mRNA /GEN=HSU	NM_015870	1.29
CDC5 cell division cycle 5-like (S. pombe)	NM_001253	1.29
zinc finger protein 16 (KOX 9)	NM_006958	1.29
alpha thalassemia/mental retardation syndrome X-linked (RAD54 homolog, S. cerevisiae)	U09820	1.29
phosphodiesterase 4D interacting protein (myomegalin)	AB042555	1.29
envoplakin	NM_001988	1.29
mitogen-activated protein kinase kinase kinase kinase 1	BE646618	1.29
ceroid-lipofuscinosis, neuronal 5	NM_006493	1.29
KIAA0507 protein	AW029169	1.29
translocated promoter region (to activated MET oncogene)	NM_003292	1.29
UDP glycosyltransferase 1 family, polypeptide A1	NM_000463	1.28
hypothetical protein FLJ20125	NM_017676	1.28
acetyl-Coenzyme A transporter	BE464756	1.28
Consensus includes gb:AI952772 /FEA=EST /DB_XREF=gi:5745082 /DB_XREF=est:wr50d05.x1 /CLONE=IMAGE:2491113 /UG=Hs.300865 Homo sapiens clone BC5ynL38 immunoglobulin l	AI952772	1.28
sema domain, immunoglobulin domain (Ig), transmembrane domain TM) and short cytoplasmic domain, (semaphorin) 4C	AI949392	1.28
pumilio (Drosophila) homolog 2	O87078	1.28
methionyl aminopeptidase 2	AI743685	1.28
LanC (bacterial lantibiotic synthetase component C)-like 1	NM_006055	1.28
CDC-like kinase 3	NM_003992	1.28
SET domain, bifurcated 1	AI762193	1.28
UDP-Gal:betaGlcNAc beta 1,4- galactosyltransferase, polypeptide 5	BF691447	1.28
Consensus includes gb:AL109658 /DEF=Human DNA sequence from clone RP4-776F14 on chromosome 20p12.2-13. Contains the 5 end of the FKBP1A gene for FK506-binding protein 1A (	AL109658	1.28
RNA binding motif protein 10	BC004181	1.28
capping protein (actin filament), gelsolin-like	NM_001747	1.28
sperm associated antigen 9	AK023512	1.28
endothelin receptor type B	M74921	1.28
hypothetical protein FLJ22679	NM_017698	1.28
GDP dissociation inhibitor 1	NM_001493	1.28
selenophosphate synthetase 2	NM_012248	1.28
checkpoint with forkhead and ring finger domains	NM_018223	1.28
adducin 3 (gamma)	NM_019903	1.28
hypothetical protein 24636	BC002640	1.28
FtsJ homolog 2 (E. coli)	NM_013393	1.28
natriuretic peptide precursor A	M30262	1.28
triple functional domain (PTPRF interacting)	AF091395	1.28
huntingtin interacting protein 1	U79734	1.28
Consensus includes gb:AV733266 /FEA=EST /DB_XREF=gi:10850811 /DB_XREF=est:AV733266 /CLONE=cdAAUG04 /UG=Hs.76325 step II splicing factor SLU7	AV733266	1.28
carbonic anhydrase II	M36532	1.28
Consensus includes gb:BG429214 /FEA=EST /DB_XREF=gi:13335720 /DB_XREF=est:602498083F1 /CLONE=IMAGE:4611836 /UG=Hs.258144 zinc finger protein 33b (KOX 31)	BG429214	1.28
heterogeneous nuclear ribonucleoprotein A3	NM_005758	1.28
KIAA1042 protein	AI633774	1.28
hypothetical protein FLJ20288	NM_024668	1.28
heparan sulfate 6-O-sulfotransferase	NM_004807	1.28
potassium voltage-gated channel, shaker-related subfamily, member 6	AW134812	1.28

tetratricopeptide repeat domain 3	AW510696	1.28
ATPase, H+ transporting, lysosomal (vacuolar proton pump) 9kD	NM_003945	1.28
IMP (inosine monophosphate) dehydrogenase 2	NM_000884	1.28
O-linked N-acetylglucosamine (GlcNAc) transferase (UDP-N-acetylglucosamine:polypeptide-N-acetylglucosaminyl transferase)	NM_003605	1.28
transporter 2, ATP-binding cassette, sub-family B (MDR/TAP)	NM_000544	1.28
arginine vasopressin receptor 1A	AI005066	1.28
putative translation initiation factor	AL523310	1.28
chondroitin sulfate proteoglycan 6 (bamacan)	BF795297	1.28
PTK6 protein tyrosine kinase 6	NM_005975	1.28
hypothetical protein MGC10763	AJ560217	1.28
hypothetical protein FLJ10241	NM_018035	1.28
KIAA1041 protein	NM_014947	1.28
calcium/calmodulin-dependent protein kinase kinase 2, beta	AA181179	1.28
IMS protein	BC000464	1.28
tryptophan rich basic protein	NM_004627	1.28
H1 histone family, member 3	NM_005320	1.28
DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 9 (RNA helicase A, nuclear DNA helicase II; leukophysin)	BF313832	1.28
KIAA1185 protein	AB033011	1.28
z 006 protein	NM_020233	1.28
ribosomal protein L22	NM_000983	1.28
H1 histone family, member 0	Z97630	1.28
6 protein-binding protein CRF6	AI081107	1.28
zinc finger protein 217	NM_006526	1.28
Consensus includes gb:BC004327.1 /DEF=Homo sapiens, clone IMAGE:3629896, mRNA, partial cds. /FEA=mRNA /PROD=Unknown (protein for IMAGE:3629896) /DB_XREF=gi:13279244 /L	BC004327	1.28
calponin 2	NM_004368	1.28
stress-associated endoplasmic reticulum protein 1; ribosome associated membrane protein 4	AL136807	1.28
hypothetical protein FLJ10900	NM_018264	1.28
NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 2 (8kD, B8)	BC003674	1.28
associated molecule with the SH3 domain of STAM	NM_006463	1.28
RAB4, member RAS oncogene family	BC002438	1.28
regulator of G-protein signalling 14	AF037194	1.28
myosin, heavy polypeptide 6, cardiac muscle, alpha (cardiomyopathy, hypertrophic 1)	D00943	1.28
splicing factor, arginine/serine-rich 2	NM_003016	1.28
Consensus includes gb:AK000145.1 /DEF=Homo sapiens cDNA FLJ20138 fis, clone COL07172. /FEA=mRNA /DB_XREF=gi:7020041 /UG=Hs.306384 Homo sapiens cDNA FLJ20138 fis, clon	AK000145	1.28
FK506 binding protein 1A (12kD)	BC005147	1.28
alpha-actinin-2-associated LIM protein	BC001017	1.28
hypothetical protein	AF091086	1.28
hypothetical protein FLJ11767	NM_024593	1.28
DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 11 (CHL1-like helicase homolog, S. cerevisiae)	AI983033	1.28
KIAA0182 protein	AA206161	1.28
hypothetical protein FLJ14393	AI823896	1.28
GABA(A) receptor-associated protein-like 2	AB030710	1.28
hypothetical protein FLJ21865	NM_022759	1.28
breast cancer anti-estrogen resistance 3	NM_003567	1.28
cytochrome P450, subfamily IIJ (arachidonic acid epoxygenase) polypeptide 2	NM_000775	1.28
calcium channel, voltage-dependent, L type, alpha 1D subunit	NM_000720	1.28
hypothetical protein FLJ14827	BE966197	1.28
follicle stimulating hormone receptor	M95489	1.28
block of proliferation 1	AK024840	1.28
ATPase, H+ transporting, lysosomal (vacuolar proton pump) 42kD	BF034973	1.28
WD repeat domain 10	NM_018262	1.28
hypothetical protein FLJ12671	AW294587	1.28
major histocompatibility complex, class I, J (pseudogene)	M80469	1.28
karyopherin alpha 1 (importin alpha 5)	BC002374	1.28
95 kDa retinoblastoma protein binding protein	NM_014771	1.28
interferon gamma receptor 1	NM_000416	1.28
secreted frizzled-related protein 1	AI332407	1.28
ribosomal protein L26	AL121871	1.28
TP53 target gene 1	BC002709	1.28
seven transmembrane protein TM7SF3	NM_016551	1.28
cleavage stimulation factor, 3' pre-RNA, subunit 2, 64kD	NM_001325	1.28
hypothetical protein FLJ12879	NM_024757	1.28
Consensus includes gb:AI972416 /FEA=EST /DB_XREF=gi:5769332 /DB_XREF=est:wr39d10.x1 /CLONE=IMAGE:2490067 /UG=Hs.24385 Human hbc647 mRNA sequence	AI972416	1.27

ribosomal protein L22	D17652	1.27
Consensus includes gb:AW452796 /FEA=EST /DB_XREF=gi:6993572 /DB_XREF=est:UI-H-BI3-ais-f-06-0-UI.s1 /CLONE=IMAGE:3069059 /UG=Hs.120769 Homo sapiens cDNA FLJ20463 fis,	AW452796	1.27
ribosomal protein L22	BG152979	1.27
DKFZP564J102 protein	AL080065	1.27
inositol 1,4,5-trisphosphate 3-kinase B	NM_002221	1.27
dihydrofolate reductase	BC000192	1.27
thioredoxin-like 2	AF118652	1.27
capillary morphogenesis protein 1	AI610355	1.27
proteasome (prosome, macropain) 26S subunit, non-ATPase, 4	AB033605	1.27
HSPC040 protein	NM_016098	1.27
phospholipase A2, group IVB (cytosolic)	AK000550	1.27
KIAA0650 protein	AA868754	1.27
ubiquitin specific protease 16	NM_006447	1.27
KIAA0826 protein	BF515124	1.27
supervillin	AW872377	1.27
SMAS	X75940	1.27
methionine-tRNA synthetase	AA621558	1.27
hypothetical protein MGC4174	NM_024319	1.27
hypothetical protein FLJ21820	AK021672	1.27
methylenetetrahydrofolate dehydrogenase (NADP+ dependent), methenyltetrahydrofolate cyclohydrolase, formyltetrahydrofolate synthetase	NM_005956	1.27
DKFZP586M1523 protein	BF063896	1.27
reticulocalbin 2, EF-hand calcium binding domain	BC004892	1.27
ribosomal protein L34	NM_000995	1.27
solute carrier family 4, sodium bicarbonate cotransporter, member 8	NM_004858	1.27
hypothetical protein MGC14797	AA628051	1.27
transcription factor-like 4	BE964655	1.27
lambda-crystallin	NM_015974	1.27
hypothetical protein FLJ10706	NM_018186	1.27
hypothetical protein FLJ13119	NM_024580	1.27
phenylethanolamine N-methyltransferase	NM_002686	1.27
TH1-like (Drosophila)	NM_016397	1.27
rhomboid, veinlet-like 1 (Drosophila)	NM_003961	1.27
sema domain, transmembrane domain (TM), and cytoplasmic domain, (semaphorin) 6C	NM_030913	1.27
hypothetical protein MGC3731	AW002578	1.27
RAB11A, member RAS oncogene family	AI215102	1.27
protein kinase, cAMP-dependent, catalytic, beta	AA130247	1.27
hypothetical protein FLJ21062	NM_024788	1.27
Consensus includes gb:AF035594.1 /DEF=Homo sapiens protein kinase C-alpha mRNA, partial 3 UTR. /FEA=mRNA /DB_XREF=gi:3168857 /UG=Hs.279856 Homo sapiens protein kinase C-a	AF035594	1.27
putative human HLA class II associated protein I	T67821	1.27
phorbol-12-myristate-13-acetate-induced protein 1	NM_021127	1.27
hypothetical protein MGC4342	NM_024329	1.27
COX15 homolog, cytochrome c oxidase assembly protein (yeast)	BC002382	1.27
HIRA interacting protein 3	NM_003609	1.27
hypothetical protein	NM_017446	1.27
protein kinase, cAMP-dependent, regulatory, type II, beta	NM_002736	1.27
ribosomal protein L26 pseudogene 1	NM_016093	1.27
tumor suppressing subtransferable candidate 4	NM_005706	1.27
ribosomal protein L28	AK025866	1.27
sorting nexin 3	BE962615	1.27
RING1 and YY1 binding protein	NM_012234	1.27
hypothetical protein FLJ13330	NM_025091	1.27
fibronectin type 3 and SPRY domain-containing protein	NM_024333	1.27
LIM domain kinase 2	NM_005569	1.27
tousled-like kinase 2	BF112255	1.27
hypothetical protein FLJ10204	NM_018024	1.27
aminopeptidase puromycin sensitive	AI132583	1.27
diphtheria toxin receptor (heparin-binding epidermal growth factor-like growth factor)	M60278	1.27
ribosomal protein L5	AL137958	1.27
latent transforming growth factor beta binding protein 1	AI986120	1.27
peroxiredoxin 2	L19185	1.27
solute carrier family 20 (phosphate transporter), member 2	NM_006749	1.27
LIM protein (similar to rat protein kinase C-binding enigma)	BG054550	1.27
cytochrome c oxidase subunit Va pseudogene 1	AF042162	1.27

hypothetical protein FLJ13855	NM_023079	1.27
WW domain binding protein 4 (formin binding protein 21)	AK000979	1.27
aldo-keto reductase family 1, member C1 (dihydrodiol dehydrogenase 1; 20-alpha (3-alpha)-hydroxysteroid dehydrogenase)	NM_001353	1.27
Consensus includes gb:AW438464 /FEA=EST /DB_XREF=gi:6973770 /DB_XREF=est:xu43g07.x1 /CLONE=IMAGE:2804508 /UG=Hs.288760 Homo sapiens cDNA FLJ12327 fis, clone MAMM/	AW438464	1.27
S-adenosylhomocysteine hydrolase-like 1	NM_006621	1.27
S-adenosylhomocysteine hydrolase-like 1	AJ589266	1.27
B-cell CLL/lymphoma 7C	NM_004765	1.27
endonuclease G-like 2	NM_005728	1.27
KIAA0494 gene product	NM_014774	1.27
hypothetical protein MDS018	NM_021823	1.27
KIAA0708 protein	AB014608	1.27
Cluster Incl. AA160496:zo73a03.s1 Homo sapiens cDNA, 3' end /clone=IMAGE-592492 /clone_end=3' /gb=AA160496 /gi=1735863 /ug=Hs.110407 /len=542	AA160496	1.27
cleavage stimulation factor, 3' pre-RNA, subunit 3, 77kD	NM_001326	1.27
phospholipase A2, group IIF	NM_022819	1.27
KIAA0089 protein	AA135522	1.27
thyroid receptor interacting protein 15	AA496247	1.27
src family associated phosphoprotein 2	NM_003930	1.27
Consensus includes gb:L24521.1 /DEF=Human transformation-related protein mRNA, 3' end. /FEA=mRNA /PROD=transformation-related protein /DB_XREF=gi:403459 /UG=Hs.300705 Hun	L24521	1.27
KIAA0794 protein	BG339560	1.27
ferrochelatase (protoporphyria)	AU152635	1.27
p53-induced protein	AK026607	1.27
casein kinase 1, alpha 1	BG534245	1.27
filamin A, alpha (actin binding protein 280)	AW051856	1.27
homeo box D3	BC005124	1.27
M-phase phosphoprotein 9	X98258	1.27
spastic paraplegia 4 (autosomal dominant; spastin)	AB029006	1.27
ubiquitin-conjugating enzyme E2C	NM_007019	1.27
protocadherin beta 17 pseudogene	AF152527	1.27
peptidase (mitochondrial processing) beta	NM_004279	1.26
Consensus includes gb:AK000270.1 /DEF=Homo sapiens cDNA FLJ20263 fis, clone COLF7804, highly similar to AJ131693 Homo sapiens mRNA for AKAP450 protein. /FEA=mRNA /DB_XREF	AK000270	1.26
serine palmitoyltransferase, long chain base subunit 2	BC005123	1.26
sel-1 suppressor of lin-12-like (C. elegans)	AB020335	1.26
CD2 antigen (cytoplasmic tail) binding protein 2	BF793888	1.26
Siah-interacting protein	AF275803	1.26
glucuronidase, beta	NM_000181	1.26
flotillin 1	AF117234	1.26
NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 6 (14kD, B14)	NM_002490	1.26
KIAA0767 protein	AB018310	1.26
mucin and cadherin-like	NM_021924	1.26
hypothetical protein FLJ12838	NM_024641	1.26
Consensus includes gb:AA133342 /FEA=EST /DB_XREF=gi:1690310 /DB_XREF=est:zn92h03.s1 /CLONE=IMAGE:565685 /UG=Hs.6700 ESTs	AA133342	1.26
hypothetical protein FLJ22655	NM_024730	1.26
fragile X mental retardation 1	AA830884	1.26
KIAA0553 protein	AU146596	1.26
general transcription factor II, i	AF035737	1.26
filamin A, alpha (actin binding protein 280)	AJ625550	1.26
hypothetical protein MGC2654	NM_024109	1.26
homologous to yeast nitrogen permease (candidate tumor suppressor)	NM_006545	1.26
dTDP-D-glucose 4,6-dehydratase	NM_014305	1.26
FH1/FH2 domain-containing protein	NM_013241	1.26
KIAA1547 protein	AW873621	1.26
biphenyl hydrolase-like (serine hydrolase; breast epithelial mucin-associated antigen)	NM_004332	1.26
T-cell leukemia/lymphoma 6	AF195821	1.26
BTG family, member 2	BG339064	1.26
presenilin 2 (Alzheimer disease 4)	U34349	1.26
yeast Sec31p homolog	AB020712	1.26
ZW10 homolog, centromere/kinetochore protein (Drosophila)	NM_004724	1.26
KIAA0630 protein	BG111260	1.26
KIAA0876 protein	BE256900	1.26
hypothetical protein FLJ10781	NM_018215	1.26
hypothetical protein FLJ21047	NM_024569	1.26
farnesyl-diphosphate farnesyltransferase 1	AA872727	1.26
HLA-B associated transcript 8	NM_006709	1.26

matrix metalloproteinase 15 (membrane-inserted)	NM_002428	1.26
DKFZP547E1010 protein	NM_015607	1.26
hypothetical protein DKFZp434L0117	NM_022778	1.26
Consensus includes gb:AK026161.1 /DEF=Homo sapiens cDNA: FLJ22508 fis, clone HRC11772. /FEA=mRNA /DB_XREF=gi:10438922 /UG=Hs.8859 Homo sapiens, Similar to RIKEN cDNA 5	AK026161	1.26
hypothetical protein dJ473B4	NM_019556	1.26
ras-like protein	BF978689	1.26
heterogeneous nuclear ribonucleoprotein H3 (2H9)	L32610	1.26
mitogen-activated protein kinase-activated protein kinase 3	NM_004635	1.26
transforming growth factor, beta 2	NM_003238	1.26
NIF3 NG61 interacting factor 3-like 1 (S. pombe)	NM_021824	1.26
DKFZP434B103 protein	AF078842	1.26
RAN binding protein 3	NM_003624	1.26
hypothetical protein FLJ20241	AW183030	1.26
hypothetical protein FLJ10349	NM_018066	1.26
ret proto-oncogene (multiple endocrine neoplasia and medullary thyroid carcinoma 1, Hirschsprung disease)	M31213	1.26
ATP-binding cassette, sub-family C (CFTR/MRP), member 9	NM_005691	1.26
hypothetical protein FLJ10276	NM_018045	1.26
653955 protein	BC002637	1.26
ubiquitin-activating enzyme E1-like	NM_003335	1.26
natural killer-tumor recognition sequence	AI688640	1.26
protein inhibitor of activated STAT protein PIASy	NM_015897	1.26
general transcription factor IIH, polypeptide 2 (44kD subunit)	AF078847	1.26
phosphatidic acid phosphatase type 2A	AF014403	1.26
synaptosomal-associated protein, 23kD	BC003686	1.26
chromosome 10 open reading frame 2	NM_021830	1.26
Consensus includes gb:AL133173 /DEF=Human DNA sequence from clone RP11-453N3 on chromosome 10 Contains the 3 end of a gene similar to ABCD1 (ATP-binding cassette, sub-family	AL133173	1.26
peroxisomal biogenesis factor 7	NM_000288	1.26
nuclear factor, interleukin 3 regulated	NM_005384	1.26
APG12 autophagy 12-like (S. cerevisiae)	NM_004707	1.26
Consensus includes gb:AK025177.1 /DEF=Homo sapiens cDNA: FLJ21524 fis, clone COL05921. /FEA=mRNA /DB_XREF=gi:10437640 /UG=Hs.306778 Homo sapiens cDNA: FLJ21524 fis, cl	AK025177	1.26
ribosomal protein L23a	U43701	1.26
chromosome 2 open reading frame 6	BC003398	1.26
stearoyl-CoA desaturase (delta-9-desaturase)	AF116616	1.26
nucleolar protein 1 (120kD)	NM_006170	1.26
syntaxin 4A (placental)	NM_004604	1.26
NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 5 (16kD, SGDH)	NM_002492	1.26
protein phosphatase 3 (formerly 2B), catalytic subunit, beta isoform (calcineurin A beta)	NM_021132	1.26
artaptin 1	AF124489	1.26
KIAA0508 protein	H92070	1.26
cytochrome c oxidase subunit VIIc	NM_001867	1.26
KIAA0252 protein	D87440	1.26
mucosa associated lymphoid tissue lymphoma translocation gene 1	AB026118	1.26
hypothetical protein MGC15396	BC005853	1.26
DKFZP727M111 protein	NM_015540	1.26
SH3 domain binding glutamic acid-rich protein like	AL515318	1.26
olfactory receptor, family 2, subfamily F, member 1	AC004853	1.26
TNF receptor-associated factor 4	NM_004295	1.26
timeless homolog (Drosophila)	NM_003920	1.26
3-hydroxyisobutyryl-Coenzyme A hydrolase	AW000964	1.26
crooked neck protein (crn)	NM_016652	1.25
transmembrane protein 4	BC001027	1.25
phosphatidylinositol-4-phosphate 5-kinase, type II, beta	BF338509	1.25
polymerase (DNA directed), alpha	NM_016937	1.25
Consensus includes gb:AK021738.1 /DEF=Homo sapiens cDNA FLJ11676 fis, clone HEMBA1004752, highly similar to Homo sapiens mRNA for LAK-4p. /FEA=mRNA /DB_XREF=gi:1043298	AK021738	1.25
Kruppel-like factor 12	N49268	1.25
hypothetical protein FLJ10546	NM_018133	1.25
Consensus includes gb:AJ302559 /DEF=Homo sapiens 6M1-4P*01 pseudogene, cell line LG2 /FEA=CDS /DB_XREF=gi:12054348 /UG=Hs.307110 Homo sapiens 6M1-4P*01 pseudogene, c	AJ302559	1.25
hypothetical protein FLJ11269	NM_018372	1.25
pyruvate dehydrogenase (lipoamide) alpha 1	BF739979	1.25
PRO2000 protein	NM_014109	1.25
tubulin-specific chaperone e	NM_003193	1.25
leucine zipper-EF-hand containing transmembrane protein 1	AI359368	1.25
estrogen receptor 2 (ER beta)	AB006590	1.25

Consensus includes gb:BF221673 /FEA=EST /DB_XREF=gi:11128850 /DB_XREF=est:7o60b03.x1 /CLONE=IMAGE:3578429 /UG=Hs.287586 Homo sapiens cDNA FLJ13648 fis, clone PLACE	BF221673	1.25
glycine C-acetyltransferase (2-amino-3-ketobutyrate coenzyme A ligase)	NM_014291	1.25
phosphoribosylformylglycinamidine synthase (FGAR amidotransferase)	AL044326	1.25
cyclin M4	NM_020184	1.25
hypothetical protein FLJ13096	NM_025000	1.25
splicing factor, arginine/serine-rich 5	U30884	1.25
DR1-associated protein 1 (negative cofactor 2 alpha)	NM_006442	1.25
hypothetical protein	NM_016399	1.25
Consensus includes gb:AK024988.1 /DEF=Homo sapiens cDNA: FLJ21335 fis, clone COL02546. /FEA=mRNA /DB_XREF=gi:10437419 /UG=Hs.303001 Homo sapiens cDNA: FLJ21335 fis, cl	AK024988	1.25
multifunctional polypeptide similar to SAICAR synthetase and AIR carboxylase	AA902652	1.25
karyopherin (importin) beta 1	BG249565	1.25
similar to yeast Upf3, variant B	NM_023010	1.25
Rho guanine nucleotide exchange factor (GEF) 1	NM_004706	1.25
endosome-associated FYVE-domain protein	NM_014733	1.25
thyroid hormone receptor interactor 7	AF274949	1.25
hypothetical protein FLJ20343	NM_017775	1.25
Consensus includes gb:AL031714 /DEF=Human DNA sequence from clone LA16-35887 on chromosome 16 Contains the UBE2I gene for ubiquitin-conjugating enzyme E2I (homologous to y	AL031714	1.25
KIAA0263 gene product	BE614199	1.25
ecotropic viral integration site 5	AF008915	1.25
origin recognition complex, subunit 4-like (yeast)	AF047598	1.25
HSPC128 protein	NM_014167	1.25
serine/threonine kinase 11 (Peutz-Jeghers syndrome)	AF035625	1.25
KIAA0475 gene product	BF115776	1.25
Consensus includes gb:AK024456.1 /DEF=Homo sapiens mRNA for FLJ00048 protein, partial cds. /FEA=mRNA /GEN=FLJ00048 /PROD=FLJ00048 protein /DB_XREF=gi:10440425 /UG=Hs.	AK024456	1.25
mitochondrial ribosomal protein S33	NM_016071	1.25
DKFZP586I2223 protein	BC002857	1.25
tropomyosin 1 (alpha)	NM_000366	1.25
H3 histone family, member A	NM_003529	1.25
ras homolog gene family, member	AW003733	1.25
phosphatidylinositol binding clathrin assembly protein	AV721177	1.25
Ras-related associated with diabetes	NM_004165	1.25
hypothetical protein HSPC177	NM_016410	1.25
hypothetical protein DKFZp564A022	NM_030954	1.25
reticulon 2	NM_005619	1.25
Consensus includes gb:AU147166 /FEA=EST /DB_XREF=gi:11008687 /DB_XREF=est:AU147166 /CLONE=HEMBB1002582 /UG=Hs.301683 Homo sapiens cDNA FLJ12091 fis, clone HEMBB	AU147166	1.25
Rab geranylgeranyltransferase, beta subunit	AA129753	1.25
prepronociceptin	NM_006228	1.25
RNA guanylyltransferase and 5'-phosphatase	AB009023	1.25
single Ig IL-1R-related molecule	NM_021805	1.25
Consensus includes gb:AL574186 /FEA=EST /DB_XREF=gi:12934150 /DB_XREF=est:AL574186 /CLONE=CS0DI039YM09 (3 prime) /UG=Hs.9739 glycerol-3-phosphate dehydrogenase 1 (:	AL574186	1.25
hypothetical protein MGC20727	N32831	1.25
putative mitochondrial space protein 32.1	AF050198	1.25
purinergic receptor P2X, ligand-gated ion channel, 2	NM_016318	1.25
hypothetical protein MGC20727	N32831	1.25
nucleolar protein (KKE/D repeat)	BE796327	1.25
hypothetical protein FLJ20422	AI660497	1.25
apolipoprotein E	NM_000041	1.25
protein phosphatase 2, regulatory subunit B (B56), gamma isoform	L42375	1.25
aminopeptidase puromycin sensitive	AW055008	1.25
hypothetical protein FLJ11848	NM_025155	1.25
CGI-115 protein	NM_016052	1.25
NRAS-related gene	NM_007158	1.25
solute carrier family 21 (organic anion transporter), member 12	NM_016354	1.25
cell division cycle 27	N21397	1.25
chromosome 1 open reading frame 28	AW192876	1.25
karyopherin alpha 1 (importin alpha 5)	AA652173	1.25
SH2 domain protein 1A, Duncan's disease (lymphoproliferative syndrome)	AF100539	1.25
hypothetical protein MGC10334	NM_030575	1.25
Clustar Incl. N32782:yw91g03.s1 Homo sapiens cDNA, 3' end /clone=IMAGE-259636 /clone_end=3' /gb=N32782 /gi=1153181 /ug=Hs.13138 /len=698	N32782	1.25
hypothetical protein MGC4614	NM_024294	1.25
Consensus includes gb:BG285011 /FEA=EST /DB_XREF=gi:13036541 /DB_XREF=est:602409278F1 /CLONE=IMAGE:4538275 /UG=Hs.12702 Homo sapiens mRNA; cDNA DKFZp586N012 (:	BG285011	1.25
hypothetical protein FLJ23251	NM_024818	1.25
major histocompatibility complex, class II, DP alpha 1	M27487	1.25

A20-binding inhibitor of NF-kappaB activation-2	NM_024309	1.25
zinc finger protein 144 (Mel-18)	AI560455	1.25
KIAA1298 protein	AI651213	1.25
polymerase (RNA) II (DNA directed) polypeptide L (7.6kD)	AA772747	1.25
IMP (inosine monophosphate) dehydrogenase 1	NM_000883	1.25
proline oxidase homolog	AA074145	1.25
transcription factor-like 4	N40555	1.25
solute carrier family 4 (anion exchanger), member 1, adaptor protein	NM_018158	1.25
serine/threonine kinase 24 (STE20 homolog, yeast)	AF083420	1.25
oxidoreductase UCPA	NM_020139	1.25
mitogen-activated protein kinase 14	NM_001315	1.25
spermidine/spermine N1-acetyltransferase	BE971383	1.25
N-acylaminoacyl-peptide hydrolase	J03068	1.25
somatostatin receptor 1	NM_001049	1.25
golgi autoantigen, golgin subfamily a, 2	NM_004486	1.25
Consensus includes gb:AI149535 /FEA=EST /DB_XREF=gi:3678004 /DB_XREF=est:qc70f11.x1 /CLONE=IMAGE:1714989 /UG=Hs.24064 Homo sapiens mRNA; cDNA DKFZp586N1323 (fron	AI149535	1.24
zinc finger protein 95 homolog (mouse)	NM_014569	1.24
DKFZP566C0424 protein	AL050028	1.24
NIMA (never in mitosis gene a)-related kinase 3	AI191920	1.24
CLIP-associating protein 1	AA176798	1.24
gb:NM_013307.1 /DEF=Homo sapiens non-functional folate binding protein (HSAF000381), mRNA. /FEA=CDS /GEN=HSAF000381 /PROD=non-functional folate binding protein /DB_XREF	NM_013307	1.24
fragile X mental retardation, autosomal homolog 1	NM_005087	1.24
translocase of inner mitochondrial membrane 8 homolog B (yeast)	NM_012459	1.24
Consensus includes gb:AF305057 /DEF=Homo sapiens RTS (RTS) gene, complete cds, alternatively spliced /FEA=mRNA_1 /DB_XREF=gi:11094017 /UG=Hs.180433 rTS beta protein	AF305057	1.24
hypothetical protein MGC5306	BC001972	1.24
cAMP responsive element binding protein 1	M34356	1.24
protein phosphatase 1, regulatory subunit 7	AI090331	1.24
ATPase, Ca++ transporting, cardiac muscle, slow twitch 2	AW190070	1.24
Ste20-related serine/threonine kinase	NM_014720	1.24
uncharacterized hematopoietic stem/progenitor cells protein MDS032	NM_018467	1.24
1-acylglycerol-3-phosphate O-acyltransferase 1 (lysophosphatidic acid acyltransferase, alpha)	U56417	1.24
myeloid cell leukemia sequence 1 (BCL2-related)	AI275690	1.24
hypothetical protein MGC955	NM_024097	1.24
farnesyl diphosphate synthase (farnesyl pyrophosphate synthetase, dimethylallyltranstransferase, geranyltranstransferase)	AL022163	1.24
I-kappa-B-interacting Ras-like protein 2	NM_017595	1.24
hypothetical protein FLJ23024	NM_024936	1.24
KIAA0191 protein	AI049962	1.24
folate receptor 3 (gamma)	NM_000804	1.24
CD59 antigen p18-20 (antigen identified by monoclonal antibodies 16.3A5, EJ16, EJ30, EL32 and G344)	AL513759	1.24
RNA 3'-terminal phosphate cyclase	NM_003729	1.24
hypothetical protein FLJ20327	NM_017767	1.24
CDC20 cell division cycle 20 homolog (S. cerevisiae)	NM_001255	1.24
exosome component Rrp46	NM_020158	1.24
KIAA0445 gene product	NM_014675	1.24
brain and reproductive organ-expressed (TNFRSF1A modulator)	U19178	1.24
parathyroid hormone-like hormone	J03580	1.24
cyclin-dependent kinase 5, regulatory subunit 1 (p35)	AL567411	1.24
RNA processing factor 1	NM_025065	1.24
polymerase (RNA) II (DNA directed) polypeptide F	BC003582	1.24
Consensus includes gb:AL567779 /FEA=EST /DB_XREF=gi:12921478 /DB_XREF=est:AL567779 /CLONE=CS00F034YB08 (3 prime) /UG=Hs.84775 Human transposon-like element mRNA	AL567779	1.24
retinoid X receptor, beta	BC001167	1.24
hypothetical protein R31240_1	AI435954	1.24
hypothetical protein DKFZp762A227	BC003163	1.24
serine/threonine-protein kinase PRP4 homolog	AB011108	1.24
protein (peptidyl-prolyl cis/trans isomerase) NIMA-interacting, 4 (parvulin)	BE797213	1.24
HSPC018 protein	NM_014027	1.24
protein (peptidyl-prolyl cis/trans isomerase) NIMA-interacting, 4 (parvulin)	BE674061	1.24
Consensus includes gb:AW469555 /FEA=EST /DB_XREF=gi:7039661 /DB_XREF=est:hd19g06.x1 /CLONE=IMAGE:2910010 /UG=Hs.313848 ESTs, Moderately similar to MGN_HUMAN MAGO	AW469555	1.24
U1-snRNP binding protein homolog (70kD)	NM_022717	1.24
ubiquitin-conjugating enzyme E2D 3 (UBC4/5 homolog, yeast)	BC003395	1.24
aldo-keto reductase family 1, member C2 (dihydrodiol dehydrogenase 2; bile acid binding protein; 3-alpha hydroxysteroid dehydrogenase, type III)	U05598	1.24
progesterone receptor membrane component 1	AL547946	1.24
uridine monophosphate synthetase (orotate phosphoribosyl transferase and orotidine-5'-decarboxylase)	D86227	1.24

Consensus includes gb:AL583123 /FEA=EST /DB_XREF=gi:12951784 /DB_XREF=est:AL583123 /CLONE=CS0DL009YN09 (3 prime) /UG=Hs.294005 Homo sapiens, clone IMAGE:3050476,	AL583123	1.24
KIAA0943 protein	NM_013325	1.24
mitochondrial ribosomal protein L20	NM_017971	1.24
SRY (sex determining region Y)-box 11	AB028641	1.24
A kinase (PRKA) anchor protein 8	AW341501	1.24
cerebellar degeneration-related protein (62kD)	AL582414	1.24
extra spindle poles, S. cerevisiae, homolog of	NM_012291	1.24
nucleotide-sugar transporter similar to C. elegans sqv-7	AJ005866	1.24
elastase 2, neutrophil	NM_001972	1.24
hypothetical protein FLJ12761	NM_024545	1.24
RNA guanylyltransferase and 5'-phosphatase	AB012142	1.24
G-rich RNA sequence binding factor 1	BF058465	1.24
v-Ki-ras2 Kirsten rat sarcoma 2 viral oncogene homolog	NM_004985	1.24
KIAA1240 protein	AB033066	1.24
G protein-coupled receptor 51	AF095723	1.24
ankyrin 3, node of Ranvier (ankyrin G)	AL136710	1.24
synapsin II	AW139618	1.24
frizzled homolog 1 (Drosophila)	NM_003505	1.24
Consensus includes gb:AI620911 /FEA=EST /DB_XREF=gi:4630037 /DB_XREF=est:tu05d12.x1 /CLONE=IMAGE:2250167 /UG=Hs.205125 ESTs	AI620911	1.24
zinc finger protein 24 (KOX 17)	BC003566	1.24
solute carrier family 22 (organic cation transporter), member 1-like	NM_002555	1.24
ATPase, vacuolar, 14 kD	NM_004231	1.24
DHHC1 protein	NM_016598	1.24
matrix metalloproteinase 11 (stromelysin 3)	NM_005940	1.24
ETAA16 protein	NM_019002	1.24
DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 16	NM_003587	1.24
sine oculis homeobox homolog 2 (Drosophila)	AF332197	1.24
Ras-related GTP-binding protein	NM_006570	1.24
sterol carrier protein 2	BC005911	1.24
TNF receptor-associated factor 4	AF082185	1.24
cytochrome b-5	NM_001914	1.24
ARP3 actin-related protein 3 homolog (yeast)	AL353587	1.24
chorionic somatomammotropin hormone 2	NM_022645	1.24
low density lipoprotein-related protein 1 (alpha-2-macroglobulin receptor)	BF304759	1.24
Consensus includes gb:BF001666 /FEA=EST /DB_XREF=gi:10701941 /DB_XREF=est:7g91d12.x1 /CLONE=IMAGE:3313847 /UG=Hs.12460 Homo sapiens clone 23870 mRNA sequence	BF001666	1.24
aminoadipate-semialdehyde dehydrogenase-phosphopantetheinyl transferase	AF151057	1.24
cadherin, EGF LAG seven-pass G-type receptor 3 (flamingo homolog, Drosophila)	AB011536	1.24
ADP-ribosylation factor related protein 1	NM_003224	1.24
protocadherin gamma subfamily A, 1	AF152318	1.24
calpain 7	BE349584	1.24
hypothetical protein MGC10471	NM_030818	1.24
eukaryotic translation initiation factor 4 gamma, 3	AI768122	1.23
thymosin, beta 4, Y chromosome	AC006371	1.23
hypothetical protein DKFZp434E2135	NM_030804	1.23
transformation/transcription domain-associated protein	NM_003496	1.23
A kinase (PRKA) anchor protein 8	AL050160	1.23
nicotinamide N-methyltransferase	NM_006169	1.23
runt-related transcription factor 1 (acute myeloid leukemia 1; aml1 oncogene)	D43968	1.23
transglutaminase 1 (K polypeptide epidermal type I, protein-glutamine-gamma-glutamyltransferase)	NM_000359	1.23
melanoma antigen, family D, 2	AF126181	1.23
endothelin receptor type A	NM_001957	1.23
POU domain, class 2, transcription factor 1	NM_002697	1.23
hypothetical protein FLJ10539	AI188402	1.23
hypothetical protein from EUROIMAGE 2021883	AA781143	1.23
killer cell lectin-like receptor subfamily A, member 1	NM_006611	1.23
Kruppel-like factor 15	NM_014079	1.23
cyclin-dependent kinase (CDC2-like) 10	NM_003674	1.23
KRAB-zinc finger protein SZF1-1	NM_016089	1.23
chorionic somatomammotropin hormone 1 (placental lactogen)	NM_022641	1.23
glucosidase I	BC002804	1.23
arachidonate 5-lipoxygenase	AI361850	1.23
hypothetical protein MGC2577	NM_031299	1.23
Consensus includes gb:AW006953 /FEA=EST /DB_XREF=gi:5855731 /DB_XREF=est:wt08d06.x1 /CLONE=IMAGE:2506859 /UG=Hs.237946 ESTs, Moderately similar to ALU7_HUMAN ALU S	AW006953	1.23

ribophorin II	AI560720	1.23
dickkopf homolog 4 (Xenopus laevis)	NM_014420	1.23
H3 histone family, member B	NM_003530	1.23
SSA protein SS-56	NM_018073	1.23
chloride channel, calcium activated, family member 3	NM_004921	1.23
hypothetical protein FLJ11026	NM_018303	1.23
ubiquitin-conjugating enzyme HBUCE1	NM_015983	1.23
UDP glycosyltransferase 2 family, polypeptide B28	AF177272	1.23
calpain 3, (p94)	AF127765	1.23
Conserved gene telomeric to alpha globin cluster	BC004185	1.23
peter pan homolog (Drosophila)	BC000535	1.23
Ras-GTPase-activating protein SH3-domain-binding protein	NM_005754	1.23
PFTAIRE protein kinase 1	AF119833	1.23
eukaryotic translation initiation factor 3, subunit 6 (48kD)	BC000734	1.23
hypothetical protein FLJ10468	BC001651	1.23
WD-repeat protein	BC001264	1.23
nuclear pore complex interacting protein	NM_006985	1.23
hypothetical protein FLJ20847	NM_017966	1.23
mitochondrial ribosomal protein L48	NM_016055	1.23
small nuclear ribonucleoprotein polypeptide N	NM_003097	1.23
SEC24 related gene family, member B (S. cerevisiae)	NM_006323	1.23
chromobox homolog 3 (HP1 gamma homolog, Drosophila)	NM_016587	1.23
up-regulated by BCG-CWS	AB040120	1.23
H3 histone family, member H	NM_003534	1.23
ATPase, H+ transporting, lysosomal (vacuolar proton pump) non-catalytic accessory protein 1A (110/116kD)	AL096733	1.23
arachidonate 5-lipoxygenase	AA995910	1.23
gb:BC001788.1 /DEF=Homo sapiens, Similar to hypothetical protein from EUROIMAGE 42353, clone MGC:2606, mRNA, complete cds. /FEA=mRNA /PROD=Similar to hypothetical protein fro	BC001788	1.23
breakpoint cluster region	U07000	1.23
hypothetical protein MGC5560	NM_019054	1.23
potassium inwardly-rectifying channel, subfamily J, member 6	N39314	1.23
KIAA1093 protein	N64803	1.23
UMP-CMP kinase	NM_016308	1.23
transcriptional regulator interacting with the PHD-bromodomain 2	NM_014755	1.23
msh homeo box homolog 2 (Drosophila)	D31771	1.23
putative L-type neutral amino acid transporter	AW000954	1.23
joined to JAZF1	AI924660	1.23
amyloid beta (A4) precursor protein-binding, family B, member 2 (Fe65-like)	U62325	1.23
H2A histone family, member M	NM_003513	1.23
regulator of nonsense transcripts 2	NM_015542	1.23
mitogen-activated protein kinase 9	U35002	1.23
zinc finger protein 134 (clone pHZ-15)	NM_003435	1.23
synaptojanin 2	AK026758	1.23
histone deacetylase 7A	NM_016596	1.23
DKFZP566E104 protein	AL110238	1.23
KIAA0625 protein	NM_015046	1.23
DKFZP566C134 protein	AF161402	1.23
stromal antigen 1	AW168948	1.23
acyl-Coenzyme A oxidase 3, pristanoyl	NM_003501	1.23
signal sequence receptor, alpha (translocon-associated protein alpha)	AW006345	1.23
chromosome 18 open reading frame 1	NM_004338	1.23
coagulation factor XII (Hageman factor)	NM_000505	1.23
hypothetical protein MGC3121	AA161130	1.23
regulator of G-protein signalling 10	W19676	1.23
ribophorin II	AL514285	1.23
hypothetical protein FLJ12438	W60802	1.23
DKFZP434N161 protein	NM_015493	1.23
tropomyosin 1 (alpha)	Z24727	1.23
GS2 gene	AI814551	1.23
Consensus includes gb:AI186739 /FEA=EST /DB_XREF=gi:3737377 /DB_XREF=est:qe79c01.x1 /CLONE=IMAGE:1745184 /UG=Hs.326416 Homo sapiens mRNA; cDNA DKFZp564H1916 (fr	AI186739	1.23
E74-like factor 1 (ets domain transcription factor)	M82882	1.23
early development regulator 1 (homolog of polyhomeotic 1)	NM_004426	1.23
KIAA0676 protein	AK001196	1.23
protein kinase, AMP-activated, beta 1 non-catalytic subunit	BC001007	1.23

hypothetical protein FLJ12436	NM_024661	1.23
transglutaminase 2 (C polypeptide, protein-glutamine-gamma-glutamyltransferase)	BC003551	1.22
vascular endothelial growth factor	H95344	1.22
casein kinase 1, gamma 2	AL530441	1.22
ARP1 actin-related protein 1 homolog A, centractin alpha (yeast)	AL532341	1.22
thyroid hormone receptor interactor 11	AF007217	1.22
putative dimethyladenosine transferase	BC002841	1.22
gb:BC001865.1 /DEF=Homo sapiens, Similar to cadherin 1, type 1, E-cadherin (epithelial), clone MGC:1151, mRNA, complete cds. /FEA=mRNA /PROD=Similar to cadherin 1, type 1, E-cadh	BC001865	1.22
calreticulin	AD000092	1.22
ribosomal protein S8	AA976278	1.22
spindle pole body protein	AJ003062	1.22
8-oxoguanine DNA glycosylase	NM_016820	1.22
KIAA0143 protein	AW470003	1.22
translocase of inner mitochondrial membrane 17 homolog B (yeast)	NM_005834	1.22
centromere protein E (312kD)	NM_001813	1.22
macrophage stimulating 1 (hepatocyte growth factor-like)	U37055	1.22
Consensus includes gb:AK024525.1 /DEF=Homo sapiens cDNA: FLJ20872 fis, clone ADKA02604. /FEA=mRNA /DB_XREF=gi:10436827 /UG=Hs.306683 Homo sapiens cDNA: FLJ20872 fis,	AK024525	1.22
ribosomal protein S7	AI970731	1.22
Smcy homolog, Y chromosome (mouse)	NM_004653	1.22
protein kinase C, iota	L18964	1.22
hypothetical protein FLJ13782	NM_024915	1.22
hypothetical protein MGC19556	AV746402	1.22
fem-1 homolog b (C. elegans)	AI799061	1.22
polymerase (RNA) III (DNA directed) (39kD)	NM_006466	1.22
natural killer-tumor recognition sequence	NM_005385	1.22
neural polypyrimidine tract binding protein	NM_021190	1.22
purinergic receptor P2X, ligand-gated ion channel, 5	U49396	1.22
hypothetical protein AF311304	NM_031214	1.22
KIAA1030 protein	AB028953	1.22
KIAA0197 protein	D83781	1.22
ribosomal protein S29	NM_001032	1.22
ADP-ribosyltransferase (NAD+; poly(ADP-ribose) polymerase)-like 2	AK001980	1.22
thyroid hormone receptor-associated protein, 95-kD subunit	AF106934	1.22
MCM3 minichromosome maintenance deficient 3 (S. cerevisiae)	NM_002388	1.22
uncharacterized hematopoietic stem/progenitor cells protein MDS032	BC006005	1.22
regulator of G-protein signalling 19	NM_005873	1.22
elongation of very long chain fatty acids (FEN1/Elo2, SUR4/Elo3, yeast)-like 1	NM_016031	1.22
trophinin	U04811	1.22
aladin	NM_015665	1.22
Nijmegen breakage syndrome 1 (nibrin)	AK001017	1.22
mitochondrial ribosomal protein S30	NM_016640	1.22
KIAA0140 gene product	NM_014661	1.22
karyopherin (importin) beta 2	NM_002270	1.22
KIAA0573 protein	BF439241	1.22
cytidine monophosphate-N-acetylneuraminic acid hydroxylase (CMP-N-acetylneuraminate monooxygenase)	NM_003570	1.22
septin 6	AW150913	1.22
bone morphogenetic protein 7 (osteogenic protein 1)	AL157414	1.22
kinesin heavy chain member 2	NM_004520	1.22
ariadne homolog, ubiquitin-conjugating enzyme E2 binding protein, 1 (Drosophila)	AI694332	1.22
ubinuclein 1	T70262	1.22
hypothetical protein MGC4276 similar to CG8198	BC002675	1.22
butyrophilin, subfamily 3, member A3	NM_006994	1.22
BRF1 homolog, subunit of RNA polymerase III transcription initiation factor IIIB (S.cerevisiae)	N91109	1.22
Consensus includes gb:AK022663.1 /DEF=Homo sapiens cDNA FLJ12601 fis, clone NT2RM4001414. /FEA=mRNA /DB_XREF=gi:10434187 /UG=Hs.288549 Homo sapiens cDNA FLJ12601 fi	AK022663	1.22
fibroblast growth factor 4 (heparin secretory transforming protein 1, Kaposi sarcoma oncogene)	BF510715	1.22
hypothetical protein FLJ20150	NM_017688	1.22
solute carrier 16 (monocarboxylic acid transporters), member 8	NM_013356	1.22
KIAA0155 gene product	NM_014633	1.22
fumarate hydratase	AA669797	1.22
BUP protein	NM_012071	1.22
RAD1 homolog (S. pombe)	AF084513	1.22
gb:NM_030807.1 /DEF=Homo sapiens glucose transporter protein 10 (GLUT10), mRNA. /FEA=mRNA /GEN=GLUT10 /PROD=glucose transporter protein 10 /DB_XREF=gi:13540597 /FL=g	NM_030807	1.22
hypothetical SBBI03 protein	AL583171	1.22

KIAA0671 gene product	AW664421	1.22
lysyl oxidase-like 2	NM_002318	1.22
putative zinc finger protein	AA524505	1.22
KIAA0807 protein	AB047005	1.22
ankyrin repeat domain 5	NM_022096	1.22
cyclin M3	NM_017623	1.22
peroxisome receptor 1	AW468717	1.22
H3 histone family, member L	NM_003537	1.22
tumor antigen SLP-8p	NM_016516	1.22
Friend leukemia virus integration 1	M93255	1.22
imidazoline receptor candidate	NM_007184	1.22
hepatic leukemia factor	M95585	1.22
hypothetical protein RG083M05.2	AI337901	1.22
nemo-like kinase	NM_016231	1.22
myeloid/lymphoid or mixed-lineage leukemia (trithorax homolog, Drosophila); translocated to, 10	AF060927	1.22
MSPC049 protein	NM_014149	1.22
KIAA1402 protein	AB037823	1.22
DEAD/H (Asp-Glu-Ala-Asp/His) box binding protein 1	N24868	1.22
Consensus includes gb:AI041204 /FEA=EST /DB_XREF=gi:3280398 /DB_XREF=est:ov77g06.x1 /CLONE=IMAGE:1643386 /UG=Hs.323748 Homo sapiens clone CDABP0086 mRNA sequence	AI041204	1.22
hypothetical protein FLJ22624	NM_024808	1.21
hypothetical protein FLJ10876	NM_018254	1.21
cyclin L ania-6a	NM_020307	1.21
microtubule-interacting protein that associates with TRAF3	AF230877	1.21
TBP-interacting protein	AB020636	1.21
baculoviral IAP repeat-containing 3	U37546	1.21
signal recognition particle 9kD	NM_003133	1.21
estrogen-related receptor alpha	NM_004451	1.21
Cluster Incl. AW001777:ws04f09.x1 Homo sapiens cDNA, 3' end /clone=IMAGE-2496233 /clone_end=3' /gb=AW001777 /gi=5848693 /ug=Hs.233641 /len=485	AW001777	1.21
IAK binding protein	AI056051	1.21
leukotriene b4 receptor (chemokine receptor-like 1)	U33448	1.21
ribonuclease P (14kD)	AW242755	1.21
erythrocyte membrane protein band 4.1 (elliptocytosis 1, RH-linked)	NM_004437	1.21
hypothetical protein DKFZp4348195	NM_031284	1.21
cyclic AMP phosphoprotein, 19 kD	AF084555	1.21
gb:NM_018069.1 /DEF=Homo sapiens hypothetical protein FLJ10352 (FLJ10352), mRNA. /FEA=mRNA /GEN=FLJ10352 /PROD=hypothetical protein FLJ10352 /DB_XREF=gi:8922371 /UG=	NM_018069	1.21
gamma-glutamyl hydrolase (conjugase, folylpolygammaglutamyl hydrolase)	NM_003878	1.21
hypothetical protein FLJ22729	NM_024683	1.21
hypothetical protein FLJ14038	NM_025072	1.21
non-metastatic cells 3, protein expressed in	NM_002513	1.21
KIAA0974 protein	AW241779	1.21
SRY (sex determining region Y)-box 4	AI989477	1.21
sirtuin silent mating type information regulation 2 homolog 5 (S. cerevisiae)	NM_012241	1.21
apolipoprotein A-IV	NM_000482	1.21
hypothetical protein FLJ14124	NM_024868	1.21
KIAA0170 gene product	NM_014641	1.21
phosphoribosyl pyrophosphate synthetase 1	BC001605	1.21
nipsnap homolog 1 (C. elegans)	AW083371	1.21
inhibitor of kappa light polypeptide gene enhancer in B-cells, kinase beta	AU153366	1.21
KIAA0648 protein	AB014548	1.21
hypothetical protein FLJ12816	NM_022060	1.21
thioredoxin	AF313911	1.21
capping protein (actin filament) muscle Z-line, beta	U03271	1.21
XPA binding protein 1; putative ATP(GTP)-binding protein	AB044661	1.21
golgi associated, gamma adaptin ear containing, ARF binding protein 2	AI799984	1.21
Consensus includes gb:BF796940 /FEA=EST /DB_XREF=gi:12101994 /DB_XREF=est:602258153F1 /CLONE=IMAGE:4341588 /UG=Hs.294100 ESTs	BF796940	1.21
galactose-4-epimerase, UDP-	NM_000403	1.21
DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 38	AF038391	1.21
brain and reproductive organ-expressed (TNFRSF1A modulator)	AL566299	1.21
putative protein kinase NY-REN-64 antigen	NM_016123	1.21
tissue inhibitor of metalloproteinase 2	NM_003255	1.21
CDC28 protein kinase 2	NM_001827	1.21
calcium/calmodulin-dependent protein kinase (CaM kinase) II gamma	AI093569	1.21
SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily c, member 2	BF663402	1.21

hypothetical protein FLJ12150	NM_024736	1.21
chromosome 1 open reading frame 22	NM_025191	1.21
KIAA0666 protein	AK021890	1.21
a disintegrin-like and metalloprotease (reprolysin type) with thrombospondin type 1 motif, 13	NM_017587	1.21
myeloid differentiation primary response gene (88)	U70451	1.21
hypothetical protein FLJ20958	NM_022102	1.21
hypothetical gene supported by AF038182; BC009203	AL565741	1.21
gamma-aminobutyric acid (GABA) A receptor, alpha 5	AF061785	1.21
Consensus includes gb:U79273.1 /DEF=Human clone 23933 mRNA sequence. /FEA=mRNA /DB_XREF=gi:1710239 /UG=Hs.239483 Human clone 23933 mRNA sequence	U79273	1.21
aminopeptidase puromycin sensitive	AK021884	1.21
hypothetical protein FLJ10422	NM_018091	1.21
acrosomal vesicle protein 1	NM_020110	1.21
GTPase regulator associated with the focal adhesion kinase pp125	Y10388	1.21
alveolar soft part sarcoma chromosome region, candidate 1	NM_024083	1.21
DKFZP434D1335 protein	BG054966	1.21
similar to rat myomegalin	NM_022359	1.21
Ca2+-promoted Ras inactivator	AI738591	1.21
HSPC133 protein	AF201938	1.21
serine protease inhibitor, Kazal type 1	NM_003122	1.21
Consensus includes gb:AU145411 /FEA=EST /DB_XREF=gi:11006932 /DB_XREF=est:AU145411 /CLONE=HEMBA1004753 /UG=Hs.274546 Homo sapiens cDNA FLJ10190 fis, clone HEMBA	AU145411	1.21
transmembrane 6 superfamily member 1	NM_023003	1.21
vesicle-associated membrane protein 4	NM_003762	1.21
origin recognition complex, subunit 2-like (yeast)	NM_006190	1.21
transcription factor-like 5 (basic helix-loop-helix)	AL035669	1.21
polymerase (DNA directed), epsilon 3 (p17 subunit)	BC004170	1.21
collagen, type VI, alpha 1	BE350145	1.21
dual specificity phosphatase 12	NM_007240	1.21
guanine nucleotide binding protein (G protein), gamma 7	NM_005145	1.21
L-3-hydroxyacyl-Coenzyme A dehydrogenase, short chain	BC000306	1.21
KIAA1058 protein	AW450751	1.21
stem cell growth factor; lymphocyte secreted C-type lectin	NM_002975	1.21
KIAA1078 protein	AB029001	1.21
hypothetical protein FLJ20397	NM_017802	1.21
HSPC056 protein	NM_014154	1.21
alpha-1-microglobulin/bikunin precursor	NM_001633	1.21
heparan sulfate proteoglycan 2 (perlecan)	AI991033	1.21
NADH dehydrogenase (ubiquinone) 1, subcomplex unknown, 1 (6kD, KFYI)	NM_002494	1.21
acylphosphatase 2, muscle type	NM_001108	1.21
serine palmitoyltransferase, long chain base subunit 1	NM_006415	1.21
sel-1 suppressor of lin-12-like (C. elegans)	NM_005065	1.21
leptin receptor gene-related protein	NM_017526	1.21
Consensus includes gb:AU151793 /FEA=EST /DB_XREF=gi:11013314 /DB_XREF=est:AU151793 /CLONE=NT2RP2006115 /UG=Hs.157078 Homo sapiens cDNA FLJ12793 fis, clone NT2RP2	AU151793	1.21
Consensus includes gb:AL034429 /DEF=Human DNA sequence from clone RP5-894K16 on chromosome 20q13.11-13.13. Contains a COX6C (cytochrome c oxidase subunit VIc) pseudogen	AL034429	1.21
hypothetical protein dJ434O14.3	NM_025228	1.21
peptide:N-glycanase similar to yeast PNG1	NM_018297	1.21
TAF9 RNA polymerase II, TATA box binding protein (TBP)-associated factor, 32 kD	NM_003187	1.21
CD59 antigen p18-20 (antigen identified by monoclonal antibodies 16.3A5, EJ16, EJ30, EL32 and G344)	BF983379	1.21
hypothetical protein FLJ20265	NM_017733	1.21
BUB3 budding uninhibited by benzimidazoles 3 homolog (yeast)	AF047473	1.21
gene near HD on 4p16.3 with homology to hypothetical S. pombe gene	R06783	1.21
aldo-keto reductase family 1, member C1 (dihydrodiol dehydrogenase 1; 20-alpha (3-alpha)-hydroxysteroid dehydrogenase)	S68290	1.21
transcription elongation factor A (SII), 1	X57198	1.21
hypothetical protein FLJ21865	AL044097	1.21
hypothetical protein FLJ10511	NM_018120	1.21
KIAA1449 protein	AB040882	1.21
DMC1 dosage suppressor of mck1 homolog, meiosis-specific homologous recombination (yeast)	NM_007068	1.21
NADPH oxidase 3	NM_015718	1.21
golgi SNAP receptor complex member 2	AF229796	1.20
acyl-Coenzyme A oxidase 1, palmitoyl	T62985	1.20
fibroblast growth factor 6	NM_020996	1.20
asparaginyl-tRNA synthetase	NM_004539	1.20
splicing factor, arginine/serine-rich 8 (suppressor-of-white-apricot homolog, Drosophila)	AI871457	1.20
ornithine aminotransferase (gyrate atrophy)	NM_000274	1.20

Consensus includes gb:AI910895 /FEA=EST /DB_XREF=gi:5630631 /DB_XREF=est:wd20h09.x1 /CLONE=IMAGE:2328737 /UG=Hs.19720 Homo sapiens mRNA full length insert cDNA clone	AI910895	1.20
inversin	AF039217	1.20
cathepsin D (lysosomal aspartyl protease)	NM_001909	1.20
Fzr1 protein	AB013463	1.20
nuclear respiratory factor 1	L22454	1.20
KIAA0144 gene product	NM_014847	1.20
butyrophilin, subfamily 2, member A1	AL121936	1.20
three prime repair exonuclease 1	AI243797	1.20
protein tyrosine phosphatase type IVA, member 2	BF795101	1.20
bone morphogenetic protein receptor, type II (serine/threonine kinase)	U20165	1.20
chymotrypsinogen B1	AW584011	1.20
mitogen-activated protein kinase-activated protein kinase 2	NM_004759	1.20
zinc finger protein 144 (Mel-18)	AI744229	1.20
butyrophilin, subfamily 3, member A2	AI991252	1.20
growth arrest and DNA-damage-inducible, alpha	NM_001924	1.20
EL6 protein	NM_018553	1.20
v-akt murine thymoma viral oncogene homolog 2	M95936	1.20
Consensus includes gb:AL137651.1 /DEF=Homo sapiens mRNA; cDNA DKFZp43400213 (from clone DKFZp43400213); partial cds. /FEA=mRNA /GEN=DKFZp43400213 /PROD=hypothetic	AL137651	1.20
caspase 3, apoptosis-related cysteine protease	NM_004346	1.20
BTB (POZ) domain containing 1	NM_025238	1.20
hypothetical protein MGC5509	NM_024093	1.20
v-maf musculoaponeurotic fibrosarcoma oncogene homolog F (avian)	AL021977	1.20
pleckstrin homology, Sec7 and coiled/coil domains, binding protein	L06633	1.20
serine (or cysteine) proteinase inhibitor, clade B (ovalbumin), member 6	BC004948	1.20
formin-binding protein 17	AU145053	1.20
hypothetical protein MGC17330	AL540260	1.20
homeo box B5	NM_002147	1.20
zinc finger RNA binding protein	BF445199	1.20
Consensus includes gb:BF445142 /FEA=EST /DB_XREF=gi:11510203 /DB_XREF=est:nad21b05.x1 /CLONE=IMAGE:3366081 /UG=Hs.797 nuclear transcription factor Y, alpha /FL=gb:NM_(	BF445142	1.20
hypothetical protein FLJ20195	NM_017706	1.20
ubiquitin specific protease	AF350251	1.20
tetratricopeptide repeat domain 3	NM_003316	1.20
butyrophilin, subfamily 2, member A1	NM_007049	1.20
hypothetical protein FLJ10618	AL049246	1.20
TBC1 (tre-2/USP6, BUB2, cdc16) domain family, member 1	AB029031	1.20
splicing factor 3b, subunit 3, 130kD	NM_012426	1.20
latent transforming growth factor beta binding protein 1	NM_000627	1.20
ATP-binding cassette, sub-family C (CFTR/MRP), member 1	AI539710	1.20
actin, alpha, cardiac muscle	NM_005159	1.20
kaptin (actin binding protein)	NM_007059	1.20
stratifin	BC000329	1.20
dual specificity phosphatase 5	U16996	1.20
hypothetical protein FLJ11159	NM_018343	1.20
core-binding factor, runt domain, alpha subunit 2; translocated to, 1; cyclin D-related	AF018283	1.20
rab6 GTPase activating protein (GAP and centrosome-associated)	AI922519	1.20
olfactory receptor, family 2, subfamily S, member 2	NM_019897	1.20
GrpE-like protein cochaperone	AI984421	1.20
hypothetical protein MGC12760	BC006312	1.20
kraken-like	AL157851	1.20
Splicing factor, arginine/serine-rich, 46kD	AI343248	1.20
nuclear cap binding protein subunit 1, 80kD	BG108193	1.20
potassium voltage-gated channel, subfamily G, member 2	NM_012283	1.20
syntaxin 3A	BE966922	1.20
hypothetical protein FLJ22035	NM_024523	1.20
putative L-type neutral amino acid transporter	AB007896	1.20
primase, polypeptide 1 (49kD)	NM_000946	1.20
gb:L34698.1 /DEF=Homo sapiens T-cell receptor alpha (TCRA) mRNA (HLA-A3, 29; B7, 44; DR 2, 7), complete cds. /FEA=mRNA /GEN=TCRA /PROD=T-cell receptor alpha /DB_XREF=gi:11(	L34698	1.20
DKFZP564C1940 protein	NM_014045	1.20
POU domain, class 2, transcription factor 2	BC006101	1.20
hypothetical protein FLJ10545	NM_018132	1.20
gb:M85256.1 /DEF=Homo sapiens immunoglobulin kappa-chain VK-1 (IgK) mRNA, complete cds. /FEA=mRNA /GEN=IgK /PROD=immunoglobulin kappa-chain VK-1 /DB_XREF=gi:186008 /(	M85256	1.20
hypothetical protein FLJ21415	NM_024738	1.20
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Consensus includes gb:AU145746 /FEA=EST /DB_XREF=gi:11007267 /DB_XREF=est:AU145746 /CLONE=HEMBA1005679 /UG=Hs.82193 esterase D formylglutathione hydrolase	AU145746	1.20
patched homolog (Drosophila)	BG054916	1.20
hypothetical protein P15-2	AF201942	1.20
hypothetical protein FLJ20188	NM_017703	1.20
Consensus includes gb:AF052091.1 /DEF=Homo sapiens clone 23578 mRNA sequence. /FEA=mRNA /DB_XREF=gi:3360397 /UG=Hs.159388 Homo sapiens clone 23578 mRNA sequence	AF052091	1.20
KIAA0830 protein	AF131747	1.20
Consensus includes gb:AI886796 /FEA=EST /DB_XREF=gi:5591960 /DB_XREF=est:wk20b07.x1 /CLONE=IMAGE:2412853 /UG=Hs.243901 Homo sapiens cDNA FLJ20738 fis, clone HEP082	AI886796	1.20
IQ motif containing GTPase activating protein 1	AI679073	1.20
solute carrier family 16 (monocarboxylic acid transporters), member 3	AA853175	1.20
Consensus includes gb:AL520908 /FEA=EST /DB_XREF=gi:12784401 /DB_XREF=est:AL520908 /CLONE=CS0DB002YF01 (3 prime) /UG=Hs.155489 NS1-associated protein 1 /FL=gb:AF1	AL520908	1.20
hypothetical protein FLJ20364	AF269167	1.20
KIAA0033 protein	H64760	1.20
VRK3 for vaccinia related kinase 3	BF062886	1.20
arginine-glutamic acid dipeptide (RE) repeats	AF016005	1.20
bromodomain adjacent to zinc finger domain, 2A	AI653126	1.20
Consensus includes gb:AI554879 /FEA=EST /DB_XREF=gi:4487242 /DB_XREF=est:te53e03.x1 /CLONE=IMAGE:2090428 /UG=Hs.81360 Homo sapiens cDNA: FLJ21927 fis, clone HEP0417	AI554879	1.20
cysteine conjugate-beta lyase; cytoplasmic (glutamine transaminase K, kynurenine aminotransferase)	NM_004059	1.20
ubiquitination factor E4B (UFD2 homolog, yeast)	NM_006048	1.20
Consensus includes gb:N36842 /FEA=EST /DB_XREF=gi:1157984 /DB_XREF=est:yy35g03.s1 /CLONE=IMAGE:273268 /UG=Hs.274412 similar to yeast Upf3, variant A	N36842	1.20
E-1 enzyme	NM_021204	1.20
microtubule-associated protein 4	M64571	1.20
Consensus includes gb:AI971258 /FEA=EST /DB_XREF=gi:5768084 /DB_XREF=est:wr27b08.x1 /CLONE=IMAGE:2488887 /UG=Hs.295923 seven in absentia (Drosophila) homolog 1	AI971258	1.20
KIAA1882 protein	H61826	1.20
hypothetical protein FLJ10618	AI694452	1.20
phospholipase C, gamma 2 (phosphatidylinositol-specific)	NM_002661	1.20
TcD37 homolog	AF123538	1.20
AD037 protein	AF260335	1.20
major histocompatibility complex, class II, DP beta 1	NM_002121	1.20
polymerase (RNA) II (DNA directed) polypeptide D	BF432147	1.20
HP1-BP74	NM_016287	1.20
bladder cancer associated protein	NM_006698	1.20
transient receptor potential channel 1	U31110	1.20
RNA, U transporter 1	NM_005701	1.20
protein phosphatase 3 (formerly 2B), catalytic subunit, gamma isoform (calcineurin A gamma)	S46622	1.20
thyroid hormone receptor, alpha (erythroblastic leukemia viral (v-erb-a) oncogene homolog, avian)	X55005	1.20
KIAA0923 protein	AW136988	1.20
solute carrier family 1 (glutamate/neutral amino acid transporter), member 4	AB026689	1.20
ribosomal protein S2	BF125158	1.19
glycogen synthase kinase 3 alpha	NM_019884	1.19
myotubularin related protein 2	U58033	1.19
lysosomal-associated membrane protein 1	NM_005561	1.19
bromodomain-containing 4	AI991631	1.19
DNA segment, single copy probe LNS-CAI/LNS-CAII (deleted in polyposis	BC000232	1.19
hypothetical protein MGC5363	NM_024064	1.19
hypothetical protein FLJ10477	NM_018105	1.19
KIAA0141 gene product	NM_014773	1.19
forkhead box D3	NM_012183	1.19
epidermal growth factor receptor pathway substrate 15	BF213575	1.19
caspase 9, apoptosis-related cysteine protease	AB015653	1.19
Consensus includes gb:AK021960.1 /DEF=Homo sapiens cDNA FLJ11898 fis, clone HEMBA1007322. /FEA=mRNA /DB_XREF=gi:10433269 /UG=Hs.313509 Homo sapiens cDNA FLJ11898 fi	AK021960	1.19
C3HC4-type zinc finger protein	NM_024835	1.19
Consensus includes gb:AA649851 /FEA=EST /DB_XREF=gi:2577179 /DB_XREF=est:ns53g06.s1 /CLONE=IMAGE:1187386 /UG=Hs.326457 ESTs	AA649851	1.19
olfactory receptor, family 2, subfamily F, member 1	NM_012369	1.19
hypothetical protein FLJ12089	NM_024552	1.19
H2A histone family, member C	NM_003509	1.19
hypothetical protein FLJ10826	NM_018233	1.19
fumarate hydratase	NM_000143	1.19
hypothetical protein FLJ10618	AI927944	1.19
H2B histone family, member A	BC000893	1.19
metallothionein 1G	NM_005950	1.19
hypothetical protein FLJ14225	NM_024874	1.19
mitogen-activated protein kinase kinase kinase 12	AW025150	1.19
tissue specific transplantation antigen P35B	U58766	1.19

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phosphatidylinositol glycan, class F	NM_002643	1.19
Cbp/p300-interacting transactivator, with Glu/Asp-rich carboxy-terminal domain, 2	AF109161	1.19
GT198, complete ORF	NM_013290	1.19
glutamine-fructose-6-phosphate transaminase 1	BE645771	1.19
hypothetical protein FLJ20644	NM_017917	1.19
chromobox homolog 6	AI458128	1.19
CDP-diacylglycerol--inositol 3-phosphatidyltransferase (phosphatidylinositol synthase)	NM_006319	1.19
cyclin-dependent kinase 9 (CDC2-related kinase)	NM_001261	1.19
capping protein (actin filament) muscle Z-line, alpha 2	AV685920	1.19
v-yes-1 Yamaguchi sarcoma viral related oncogene homolog	NM_002350	1.19
hypothetical protein FLJ20130	NM_017681	1.19
transcription factor-like 5 (basic helix-loop-helix)	NM_006602	1.19
N-ethylmaleimide-sensitive factor attachment protein, alpha	BC001165	1.19
protease, serine, 2 (trypsin 2)	NM_002770	1.19
hypothetical protein FLJ10751	NM_018205	1.19
replication initiation region protein (60kD)	NM_014374	1.19
VAMP (vesicle-associated membrane protein)-associated protein A (33kD)	AK000918	1.19
activated RNA polymerase II transcription cofactor 4	NM_006713	1.19
SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily a, member 2	NM_003070	1.19
LRP16 protein	NM_014067	1.19
KIAA0423 protein	AI823592	1.19
karyopherin (importin) beta 2	AI307759	1.19
solute carrier family 3 (cystine, dibasic and neutral amino acid transporters, activator of cystine, dibasic and neutral amino acid transport), member 1	M95548	1.19
adenylyl cyclase-associated protein	AA806142	1.19
ceroid-lipofuscinosis, neuronal 2, late infantile (Jansky-Bielschowsky disease)	AA602532	1.19
Consensus includes gb:BE967207 /FEA=EST /DB_XREF=gi:11773627 /DB_XREF=est:601661094R1 /CLONE=IMAGE:3916174 /UG=Hs.165590 ribosomal protein S13	BE967207	1.19
fem-1 homolog b (C. elegans)	AW139179	1.19
Consensus includes gb:AV728526 /FEA=EST /DB_XREF=gi:10837947 /DB_XREF=est:AV728526 /CLONE=HTCBBD04 /UG=Hs.62264 KIAA0937 protein	AV728526	1.19
Consensus includes gb:AF070537.1 /DEF=Homo sapiens clone 24606 mRNA sequence. /FEA=mRNA /DB_XREF=gi:3387894 /UG=Hs.17481 Homo sapiens clone 24606 mRNA sequence	AF070537	1.19
glycosylphosphatidylinositol specific phospholipase D1	AV699786	1.19
gb:NM_014925.1 /DEF=Homo sapiens KIAA1002 protein (KIAA1002), mRNA. /FEA=mRNA /GEN=KIAA1002 /PROD=KIAA1002 protein /DB_XREF=gi:7662441 /UG=Hs.102483 KIAA1002 pn	NM_014925	1.19
TcD37 homolog	AI762105	1.19
replication protein A3 (14kD)	BC005264	1.19
placental protein 13-like protein	NM_020129	1.19
Consensus includes gb:AL050035.1 /DEF=Homo sapiens mRNA; cDNA DKFZp566H0124 (from clone DKFZp566H0124). /FEA=mRNA /DB_XREF=gi:4884276 /UG=Hs.133130 Homo sapiens	AL050035	1.19
hypothetical protein FLJ13611	NM_024941	1.19
retinoic acid induced 16	NM_022749	1.19
calbindin 1, (28kD)	AF049895	1.19
thymine-DNA glycosylase	NM_003211	1.19
Consensus includes gb:AI884867 /FEA=EST /DB_XREF=gi:5590031 /DB_XREF=est:wi85g03.x1 /CLONE=IMAGE:2431732 /UG=Hs.91379 ribosomal protein L26	AI884867	1.19
gb:NM_005703.2 /DEF=Homo sapiens upstream regulatory element binding protein 1 (UREB1), mRNA. /FEA=CDS /GEN=UREB1 /PROD=upstream regulatory element binding protein 1 /DB	NM_005703	1.19
hypothetical protein FLJ22709	NM_024578	1.19
pinin, desmosome associated protein	BF508848	1.19
KIAA0370 protein	BE966299	1.19
Hpall tiny fragments locus 9C	NM_022727	1.19
SHC (Src homology 2 domain-containing) transforming protein 1	AI809967	1.19
glycogen synthase kinase 3 alpha	L40027	1.19
KIAA0332 protein	AI184562	1.19
syntaxin binding protein 2	AB002559	1.19
baculoviral IAP repeat-containing 4	U32974	1.19
fucosyltransferase 7 (alpha (1,3) fucosyltransferase)	U11282	1.19
retinoblastoma binding protein 4	X71810	1.19
KIAA0808 gene product	NM_014729	1.19
pM5 protein	AL512687	1.19
PR domain containing 10	NM_020228	1.19
Tat interaction protein 1	AK001327	1.19
HSPC134 protein	NM_014169	1.19
secreted protein of unknown function	NM_013349	1.19
pericentrin 2 (kendrin)	NM_006031	1.19
protein kinase, lysine deficient 1	AB002342	1.19
calumenin	NM_001219	1.19
type 1 tumor necrosis factor receptor shedding aminopeptidase regulator	AF106037	1.19
polymyositis/scleroderma autoantigen 1 (75kD)	NM_005033	1.19

mitochondrial ribosomal protein S4	AL136913	1.19
Consensus includes gb:AF119846.1 /DEF=Homo sapiens PRO1474 mRNA, complete cds. /FEA=mRNA /PROD=PRO1474 /DB_XREF=gi:7770128 /UG=Hs.324925 LATS (large tumor suppres	AF119846	1.19
protocadherin gamma subfamily C, 3	NM_002588	1.19
6-2 and S-phase expressed 1	BF973178	1.19
cytosolic ovarian carcinoma antigen 1	AK000353	1.19
topoisomerase (DNA) II alpha (170kD)	AL561834	1.19
ring finger protein 24	BC000213	1.19
CD81 antigen (target of antiproliferative antibody 1)	NM_004356	1.19
ROD1 regulator of differentiation 1 (S. pombe)	AW190873	1.19
coatomer protein complex, subunit beta	NM_016451	1.19
ribosomal protein S20	BF184532	1.19
Cluster Incl. W07773:zb03g04.r1 Homo sapiens cDNA, 5' end /clone=IMAGE-301014 /clone_end=5' /gb=W07773 /gi=1281785 /ug=Hs.9739 /len=533	W07773	1.19
protocadherin beta 13	NM_018933	1.19
KIAA0774 protein	AI818409	1.19
branched chain keto acid dehydrogenase E1, beta polypeptide (maple syrup urine disease)	AL531533	1.19
ribosomal protein L5	BF214492	1.19
sigma receptor (SR31747 binding protein 1)	NM_005866	1.19
electron-transferring-flavoprotein dehydrogenase	S69232	1.18
hypothetical protein FLJ20604	NM_017897	1.18
Consensus includes gb:AL050364.1 /DEF=Homo sapiens mRNA; cDNA DKFZp564G052 (from clone DKFZp564G052). /FEA=mRNA /DB_XREF=gi:4914598 /UG=Hs.302761 Homo sapiens mI	AL050364	1.18
thyroid hormone receptor-associated protein, 95-kD subunit	AW262690	1.18
cell division cycle 2-like 5 (cholinesterase-related cell division controller)	BE675449	1.18
cytochrome c oxidase subunit VIc	NM_004374	1.18
general transcription factor IIIA	BE542815	1.18
trophinin	AF349719	1.18
electron-transferring-flavoprotein dehydrogenase	NM_004453	1.18
protein tyrosine phosphatase type IVA, member 3	NM_007079	1.18
LPS-induced TNF-alpha factor	NM_004862	1.18
tryptophanyl tRNA synthetase 2 (mitochondrial)	NM_015836	1.18
histone deacetylase 5	NM_005474	1.18
transcription elongation factor B (SIII), polypeptide 1-like	NM_003197	1.18
TNF-induced protein	NM_014350	1.18
casein kinase 1, alpha 1	AF119911	1.18
ligase III, DNA, ATP-dependent	NM_002311	1.18
deleted in lymphocytic leukemia, 1	NM_005887	1.18
gb:AF271775.1 /DEF=Homo sapiens DC49 mRNA, complete cds. /FEA=mRNA /PROD=DC49 /DB_XREF=gi:12006206 /UG=Hs.307093 Homo sapiens DC49 mRNA, complete cds /FL=gb:AF2	AF271775	1.18
Consensus includes gb:AK026867.1 /DEF=Homo sapiens cDNA: FLJ23214 fis, clone ADSU01585. /FEA=mRNA /DB_XREF=gi:10439827 /UG=Hs.306891 Homo sapiens cDNA: FLJ23214 fis,	AK026867	1.18
solute carrier family 12 (potassium/chloride transporters), member 7	NM_006598	1.18
protocadherin gamma subfamily C, 3	BC006439	1.18
phosphodiesterase 4A, cAMP-specific (phosphodiesterase E2 dunce homolog, Drosophila)	U18088	1.18
lacrimal proline rich protein	BF665706	1.18
solute carrier family 7 (cationic amino acid transporter, y+ system), member 8	AL365343	1.18
A kinase (PRKA) anchor protein 11	NM_016248	1.18
KIAA0096 protein	AI806207	1.18
Consensus includes gb:AI291720 /FEA=EST /DB_XREF=gi:3934494 /DB_XREF=est:qm80c06.x1 /CLONE=IMAGE:1895050 /UG=Hs.262642 ESTs	AI291720	1.18
KIAA0441 gene product	NM_014797	1.18
low density lipoprotein receptor (familial hypercholesterolemia)	AI861942	1.18
solute carrier family 26, member 6	AF288410	1.18
chorionic somatomammotropin hormone 1 (placental lactogen)	NM_022642	1.18
ring finger protein 1	Z14000	1.18
KIAA0304 gene product	AF105279	1.18
ring finger protein 10	AL578551	1.18
fatty-acid-Coenzyme A ligase, long-chain 3	AL525798	1.18
CD27-binding (Siva) protein	AW024335	1.18
hypothetical protein MGC3079	BC001030	1.18
sulfotransferase family, cytosolic, 1A, phenol-preferring, member 1	U37025	1.18
SREBP CLEAVAGE-ACTIVATING PROTEIN	D83782	1.18
Cluster Incl. AI962879:wt24c06.x1 Homo sapiens cDNA, 3' end /clone=IMAGE-2508394 /clone_end=3' /gb=AI962879 /gi=5755592 /ug=Hs.81920 /len=490	AI962879	1.18
ribosomal protein L7	AL031577	1.18
C9orf10 protein	NM_014612	1.18
methylmalonate-semialdehyde dehydrogenase	AW612403	1.18
gb:BC000110.1 /DEF=Homo sapiens, Similar to E2F transcription factor 5, clone MGC:1262, mRNA, complete cds. /FEA=mRNA /PROD=Similar to E2F transcription factor 5 /DB_XREF=gi:12	BC000110	1.18
NADH dehydrogenase (ubiquinone) 1, alpha/beta subcomplex, 1 (8kD, SDAP)	NM_005003	1.18

centrin, EF-hand protein, 2	BC005334	1.18
fibroblast growth factor 12B	NM_004113	1.18
triple functional domain (PTPRF interacting)	BF223718	1.18
novel RGD-containing protein	NM_006571	1.18
nuclear antigen Sp100	NM_003113	1.18
nuclear receptor co-repressor 1	NM_006311	1.18
chromosome 22 open reading frame 5	NM_012264	1.18
Consensus includes gb:AI721229 /FEA=EST /DB_XREF=gi:5038485 /DB_XREF=est:as68c10.x1 /CLONE=IMAGE:2333874 /UG=Hs.326456 hypothetical protein FLJ20030	AI721229	1.18
F-box only protein 7	NM_012179	1.18
vesicle docking protein p115	BE875592	1.18
amyloid beta (A4) precursor protein-binding, family B, member 2 (Fe65-like)	U62325	1.18
hypothetical protein	AA748492	1.18
islet cell autoantigen 1 (69kD)	NM_004968	1.18
KIAA0342 gene product	AA035414	1.18
zinc finger protein, subfamily 1A, 1 (Ikaros)	NM_006060	1.18
hypothetical protein FLJ13910	H65865	1.18
protein z 0001	NM_016302	1.18
growth suppressor 1	NM_022356	1.18
hypothetical protein FLJ23182	NM_022366	1.18
Mid-1-related chloride channel 1	AW195581	1.18
hypothetical protein FLJ20296	NM_017750	1.18
polybromo 1	NM_018165	1.18
hypothetical protein FLJ11209	NM_018360	1.18
arrestin, beta 1	NM_004041	1.18
hypothetical protein MGC2821	AA402435	1.18
suppressor of variegation 3-9 homolog 1 (Drosophila)	NM_003173	1.18
aldehyde dehydrogenase 4 family, member A1	NM_003748	1.18
tetratricopeptide repeat domain 4	W22690	1.18
HMT1 hnRNP methyltransferase-like 1 (S. cerevisiae)	AL570294	1.18
Consensus includes gb:AI671747 /FEA=EST /DB_XREF=gi:4851478 /DB_XREF=est:wa05e06.x1 /CLONE=IMAGE:2297218 /UG=Hs.81360 Homo sapiens cDNA: FLJ21927 fis, clone HEP041	AI671747	1.18
SH2 domain protein 1A, Duncan's disease (lymphoproliferative syndrome)	AF100540	1.18
chromosome 18 open reading frame 1	AI349506	1.18
Consensus includes gb:AK025152.1 /DEF=Homo sapiens cDNA: FLJ21499 fis, clone COL05634. /FEA=mRNA /DB_XREF=gi:10437612 /UG=Hs.306773 Homo sapiens cDNA: FLJ21499 fis, cl	AK025152	1.18
UDP-N-acetylglucosamine-2-epimerase/N-acetylmannosamine kinase	NM_005476	1.18
v-raf-1 murine leukemia viral oncogene homolog 1	X06409	1.18
nicotinamide nucleotide transhydrogenase	U40490	1.18
mannosidase, alpha, class 1B, member 1	AA631254	1.18
Consensus includes gb:BF436315 /FEA=EST /DB_XREF=gi:11448630 /DB_XREF=est:7p06b05.x1 /CLONE=IMAGE:3644888 /UG=Hs.296338 ESTs	BF436315	1.18
hypothetical protein FLJ22672	NM_024897	1.18
microfibrillar-associated protein 3	BE222709	1.18
dynein intermediate chain 2	NM_023036	1.18
DKFZP564K247 protein	BE739519	1.18
transcriptional adaptor 3 (ADA3, yeast homolog)-like (PCAF histone acetylase complex)	R59977	1.18
kinesin family member 4A	NM_012310	1.18
hypothetical protein MGC5139	BC004815	1.18
sulfotransferase family, cytosolic, 1A, phenol-preferring, member 2	U28169	1.18
myelin protein zero-like 1	AF239756	1.18
mitochondrial translational initiation factor 2	NM_002453	1.18
KIAA0916 protein	AA488899	1.18
myeloid leukemia factor 2	NM_005439	1.18
selenoprotein X, 1	NM_016332	1.18
CGI-90 protein	NM_016033	1.18
neurogranin (protein kinase C substrate, RC3)	NM_006176	1.18
potassium channel, subfamily K, member 1 (TWIK-1)	NM_002245	1.18
ribosomal protein L23a	NM_000984	1.18
hypothetical protein FLJ13111	NM_025082	1.18
GTP binding protein 1	U87964	1.18
FSH primary response (LRPR1 homolog, rat) 1	NM_006733	1.18
homolog of mouse quaking QKI (KH domain RNA binding protein)	AF142421	1.18
dynamin 1	AF035321	1.18
hypothetical protein FLJ13164	NM_022776	1.18
diacylglycerol kinase, delta (130kD)	NM_003648	1.18
cat eye syndrome chromosome region, candidate 1	NM_017424	1.18

inositol polyphosphate-5-phosphatase, 75kD	AK022846	1.18
chromosome 9 open reading frame 7	AL042496	1.18
inositol 1,4,5-triphosphate receptor, type 2	NM_002223	1.18
kynurenine 3-monooxygenase (kynurenine 3-hydroxylase)	NM_003679	1.18
male-specific lethal 3-like 1 (Drosophila)	R10150	1.18
tubulin-specific chaperone a	NM_004607	1.18
spermidine/spermine N1-acetyltransferase	M55580	1.18
KIAA0164 gene product	NM_014739	1.18
hypothetical protein R30953_1	AA431984	1.18
hypothetical protein FLJ20808	NM_017960	1.18
Consensus includes gb:AL359954 /DEF=Human DNA sequence from clone RP11-318P23 on chromosome 20 Contains a TAR DNA-binding protein pseudogene, ESTs, STSs and GSSs /FEA=CI	AL359954	1.18
integrin beta 1 binding protein (melusin) 2	NM_012278	1.17
6 protein-coupled receptor kinase-interactor 2	AF124491	1.17
transcription factor A, mitochondrial	NM_003201	1.17
leucine zipper, putative tumor suppressor 1	AF123652	1.17
phosphatidylinositol-3 phosphate 3-phosphatase adaptor subunit	NM_019061	1.17
cystatin S	NM_001899	1.17
Consensus includes gb:Z98751 /DEF=Human DNA sequence from PAC 56089 on chromosome 1q24-1q25. Contains profilin-like pseudogene, 60S ribosomal protein L4 pseudogene RNA bin	Z98751	1.17
suppression of tumorigenicity 13 (colon carcinoma) (Hsp70 interacting protein)	U17714	1.17
hypothetical protein FLJ10560	NM_018138	1.17
immunoglobulin kappa variable 1/OR1S-118	M20812	1.17
carnitine palmitoyltransferase II	NM_000098	1.17
RecQ protein-like 5	AB006533	1.17
heat shock 105kD	BG403660	1.17
hypothetical protein FLJ21106	NM_025097	1.17
hypothetical protein PRO2389	AF283773	1.17
butyrophilin, subfamily 2, member A2	W58757	1.17
general transcription factor IIIC, polypeptide 2 (beta subunit, 110kD)	BC000212	1.17
enigma (LIM domain protein)	AA086229	1.17
protein tyrosine phosphatase, receptor type, f polypeptide (PTPRF), interacting protein (liprin), alpha 1	U22815	1.17
Consensus includes gb:BE503392 /FEA=EST /DB_XREF=gi:9705800 /DB_XREF=est:7a17e04.x1 /CLONE=IMAGE:3219006 /UG=Hs.145020 ESTs, Weakly similar to KIAA1205 protein H.sapi	BE503392	1.17
protection of telomeres 1	NM_015450	1.17
N-acylsphingosine amidohydrolase (acid ceramidase)-like	AV724215	1.17
caspase recruitment domain protein 10	AY028896	1.17
Glutamate dehydrogenase-2	AC006144	1.17
cold shock domain protein A	AL556190	1.17
integral membrane protein 2B	AF092128	1.17
TEA domain family member 4	U63824	1.17
spondyloepiphyseal dysplasia, late	NM_014563	1.17
CD36 antigen (collagen type I receptor, thrombospondin receptor)-like 1	NM_005505	1.17
hypothetical protein MGC5347	NM_024063	1.17
protein tyrosine kinase 9	NM_002822	1.17
selectin P ligand	U02297	1.17
Consensus includes gb:AL049842 /DEF=Human DNA sequence from clone 129L7 on chromosome 6q12-13. Contains the gene for a PUTATIVE novel protein, ESTs, an STS, GSSs and a taga n	AL049842	1.17
transmembrane protein 1	U61500	1.17
MpV17 transgene, murine homolog, glomerulosclerosis	NM_002437	1.17
testis specific ankyrin-like protein 1	NM_016552	1.17
SH3 protein interacting with Nck, 90 kDa	NM_016453	1.17
HLA class II region expressed gene KE4	NM_006979	1.17
GAP-associated tyrosine phosphoprotein p62 (Sam68)	NM_006559	1.17
hypothetical protein FLJ20533	NM_017866	1.17
DKFZP547E1010 protein	AW005776	1.17
KIAA1449 protein	AI806628	1.17
Consensus includes gb:AV728268 /FEA=EST /DB_XREF=gi:10837689 /DB_XREF=est:AV728268 /CLONE=HTCACH05 /UG=Hs.82845 Homo sapiens cDNA: FLJ21930 fis, clone HEP04301, hi	AV728268	1.17
BTAF1 RNA polymerase II, B-TFIID transcription factor-associated, 170 kD (Mot1 homolog, S. cerevisiae)	AI001017	1.17
pumilio homolog 1 (Drosophila)	NM_014676	1.17
hemoglobin, epsilon 1	NM_005330	1.17
nucleolar and coiled-body phosphprotein 1	AI355279	1.17
signal transducer and activator of transcription 3 (acute-phase response factor)	BC000627	1.17
putative Rab5-interacting protein	NM_018840	1.17
protein phosphatase 6, catalytic subunit	NM_016294	1.17
KIAA0152 gene product	BC000371	1.17
hypothetical protein FLJ10955	NM_018282	1.17

CGI-87 protein	NM_016030	1.17
tropomodulin	BC002660	1.17
splicing factor 3b, subunit 1, 155kD	AW003030	1.17
thyroid hormone receptor-associated protein, 95-kD subunit	NM_005481	1.17
actin, gamma 2, smooth muscle, enteric	NM_001615	1.17
hypothetical protein FLJ12270	NM_024673	1.17
casein kinase 1, epsilon	NM_001894	1.17
inositol polyphosphate-5-phosphatase, 145kD	U53470	1.17
protein kinase, X-linked	NM_005044	1.17
solute carrier family 15 (H+/peptide transporter), member 2	NM_021082	1.17
core promoter element binding protein	BE675435	1.17
Consensus includes gb:S81916.1 /DEF=phosphoglycerate kinase (alternatively spliced) human, phosphoglycerate kinase deficient patient with episodes of muscl, mRNA Partial Mutant, 307 r	S81916	1.17
fragile X mental retardation 2	NM_002025	1.17
tubby like protein 3	AJ591305	1.17
SEC14-like 1 (S. cerevisiae)	AV748469	1.17
Pseudoautosomal GTP-binding protein-like	NM_012227	1.17
hypothetical protein FLJ10867	NM_018249	1.17
hypothetical protein FLJ20272	NM_017735	1.17
cullin 4B	AV694732	1.17
mitochondrial ribosomal protein S14	BG254653	1.17
WW domain-containing protein 1	BF131791	1.17
RAN binding protein 2	AI681120	1.17
zinc finger protein 225	NM_013362	1.17
hypothetical protein FLJ22471	NM_025140	1.17
KIAA0185 protein	AW026194	1.17
hypothetical protein FLJ21959	NM_022917	1.17
Consensus includes gb:AK000834.1 /DEF=Homo sapiens cDNA FLJ20827 fis, clone ADKA03543. /FEA=mRNA /DB_XREF=gi:7021152 /UG=Hs.306409 Homo sapiens cDNA FLJ20827 fis, clo	AK000834	1.17
G protein-coupled receptor kinase 6	AF040752	1.17
lipoyltransferase	NM_015929	1.17
zinc finger protein 207	NM_003457	1.17
olfactory receptor, family 2, subfamily A, member 9 pseudogene	AA731709	1.17
growth arrest and DNA-damage-inducible, beta	AF087853	1.17
calcium binding atopy-related autoantigen 1	AK022697	1.17
tumor protein p53 binding protein, 1	NM_005657	1.17
kinesin family member 5B	NM_004521	1.17
retinoic acid repressible protein	AF130102	1.17
COX11 homolog, cytochrome c oxidase assembly protein (yeast)	AJ376724	1.17
primase, polypeptide 2A (58kD)	NM_000947	1.17
a disintegrin and metalloproteinase domain 23	NM_003812	1.17
protein tyrosine phosphatase, non-receptor type 7	NM_002832	1.17
plectin 1, intermediate filament binding protein, 500kD	Z54367	1.17
interleukin 2 receptor, gamma (severe combined immunodeficiency)	NM_000206	1.17
regulatory solute carrier protein, family 1, member 1	AI268381	1.17
proliferation-associated 2G4, 38kD	BE221922	1.17
hypothetical protein FLJ10074	NM_017988	1.17
gb:AF249277.1 /DEF=Homo sapiens cervical cancer suppressor-1 mRNA, complete cds. /FEA=mRNA /PROD=cervical cancer suppressor-1 /DB_XREF=gi:10442005 /UG=Hs.314544 Homo s	AF249277	1.17
nuclear transcription factor Y, alpha	NM_002505	1.17
potassium intermediate/small conductance calcium-activated channel, subfamily N, member 1	NM_002248	1.17
calmodulin 2 (phosphorylase kinase, delta)	NM_001743	1.17
Fanconi anemia, complementation group E	NM_021922	1.17
KIAA0391 gene product	AA129755	1.17
nucleosome assembly protein 1-like 1	AI888672	1.17
calcium channel, voltage-dependent, alpha 1I subunit	AF211189	1.16
endothelial differentiation, lysophosphatidic acid G-protein-coupled receptor, 7	NM_012152	1.16
vesicle-associated membrane protein 2 (synaptobrevin 2)	BC002737	1.16
hypothetical protein FLJ20294	NM_017749	1.16
ribosome binding protein 1 homolog 180kD (dog)	NM_004587	1.16
BTB and CNC homology 1, basic leucine zipper transcription factor 2	NM_021813	1.16
gamma-glutamyltransferase-like 3	AL049709	1.16
nuclear receptor subfamily 1, group D, member 2	N32859	1.16
mitochondrial ribosomal protein S35	NM_021821	1.16
mitogen-activated protein kinase kinase kinase kinase 1	NM_007181	1.16
ubiquitin specific protease 11	BC000350	1.16

signal transducer and activator of transcription 1, 91kD	M97935	1.16
cortactin SH3 domain-binding protein	AF131790	1.16
hemoglobin, gamma A	NM_000559	1.16
golgin-67	AF204231	1.16
microtubule-associated proteins 1A/1B light chain 3	BE893893	1.16
G protein-coupled receptor 2	NM_016602	1.16
regulator of G-protein signalling 11	NM_003834	1.16
cleft lip and palate associated transmembrane protein 1	NM_001294	1.16
KIAA0876 protein	BE256900	1.16
muscleblind-like (Drosophila)	NM_021038	1.16
KIAA0118 protein	NM_014999	1.16
hypothetical protein FLJ20396	NM_017801	1.16
hypothetical protein RG083M05.2	AC000064	1.16
solute carrier family 31 (copper transporters), member 1	NM_001859	1.16
PC4 and SFRS1 interacting protein 2	AF063020	1.16
flotillin 2	NM_004475	1.16
retinal degeneration B beta	NM_012417	1.16
hypothetical protein FLJ21156	NM_024602	1.16
egl nine homolog 2 (C. elegans)	NM_017555	1.16
sequestosome 1	NM_003900	1.16
gb:NM_020412.1 /DEF=Homo sapiens CHMP1.5 protein (CHMP1.5), mRNA. /FEA=mRNA /GEN=CHMP1.5 /PROD=CHMP1.5 protein /DB_XREF=gi:9966900 /UG=Hs.42733 CHMP1.5 protein	NM_020412	1.16
cell matrix adhesion regulator	NM_005200	1.16
NCK-associated protein 1	NM_013436	1.16
PTD015 protein	BC002752	1.16
MLL septin-like fusion	AF142408	1.16
PDZ domain containing guanine nucleotide exchange factor(GEF)1	NM_014247	1.16
chromodomain protein, Y chromosome-like	AL050164	1.16
phosphatidylinositol glycan, class C, pseudogene 1	AB000359	1.16
Consensus includes gb:AA769006 /FEA=EST /DB_XREF=gi:2820244 /DB_XREF=est:oa80c10.s1 /CLONE=IMAGE:1318578 /UG=Hs.290921 ESTs, Highly similar to G02750 DNA-directed DN	AA769006	1.16
beclin 1 (coiled-coil, myosin-like BCL2 interacting protein)	NM_003766	1.16
RAN binding protein 9	AA191426	1.16
sin3-associated polypeptide, 18kD	BF593650	1.16
gb:NM_001119.2 /DEF=Homo sapiens adducin 1 (alpha) (ADD1), transcript variant 1, mRNA. /FEA=mRNA /GEN=ADD1 /PROD=adducin 1 (alpha) isoform a /DB_XREF=gi:7710113 /FL=gb	NM_001119	1.16
HSPC052 protein	NM_014150	1.16
ceroid-lipofuscinosis, neuronal 3, juvenile (Batten, Spielmeyer-Vogt disease)	AF015593	1.16
intercellular adhesion molecule 5, telencephalin	NM_003259	1.16
hypothetical protein FLJ11413	NM_024554	1.16
WD repeat domain 3	NM_006784	1.16
Consensus includes gb:BE673445 /FEA=EST /DB_XREF=gi:10033986 /DB_XREF=est:7d35d10.x1 /CLONE=IMAGE:3249235 /UG=Hs.22049 Homo sapiens chromosome 19, cosmid R28379	BE673445	1.16
mitochondrial ribosomal protein S12	AAS13737	1.16
DKFZP564K196 protein	AL049329	1.16
cytochrome c oxidase subunit VIc pseudogene 1	AF042163	1.16
leucine-rich repeat-containing 2	NM_024512	1.16
hypothetical protein FLJ21047	BF978611	1.16
spectrin, beta, non-erythrocytic 2	NM_006946	1.16
zinc finger protein	NM_014347	1.16
mucin 8, tracheobronchial	U14383	1.16
butyrophilin, subfamily 3, member A3	NM_006994	1.16
IKK-related kinase epsilon; inducible IkappaB kinase	AW340333	1.16
likely homolog of rat kinase D-interacting substance of 220 kDa	AK022873	1.16
polymerase (RNA) II (DNA directed) polypeptide A (220kD)	NM_000937	1.16
transcription termination factor, RNA polymerase II	AF080255	1.16
v-myc myelocytomatosis viral related oncogene, neuroblastoma derived (avian)	AI871354	1.16
BBP-like protein 2	AF353992	1.16
calreticulin	BE251303	1.16
MLL septin-like fusion	NM_006640	1.16
immunoglobulin kappa variable 1/OR2-108	X51887	1.16
Consensus includes gb:AK024252.1 /DEF=Homo sapiens cDNA FLJ14190 fis, clone NT2RP2006534, moderately similar to 5-AMP-ACTIVATED PROTEIN KINASE, CATALYTIC ALPHA-1 CHAIN (EC	AK024252	1.16
ATP-binding cassette, sub-family C (CFTR/MRP), member 8	AF087138	1.16
DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 24	NM_020414	1.16
gb:NM_025156.1 /DEF=Homo sapiens hypothetical protein FLJ12474 (FLJ12474), mRNA. /FEA=mRNA /GEN=FLJ12474 /PROD=hypothetical protein FLJ12474 /DB_XREF=gi:13376752 /UG	NM_025156	1.16
splicing factor, arginine/serine-rich 8 (suppressor-of-white-apricot homolog, Drosophila)	AI023864	1.16
spindle pole body protein	NM_006322	1.16

makorin, ring finger protein, 1	NM_013446	1.16
golgi autoantigen, golgin subfamily a, 1	NM_002077	1.16
hypothetical protein FLJ20311	NM_017760	1.16
DKFZP564M182 protein	AK025446	1.16
MHC class I polypeptide-related sequence B	NM_005931	1.16
chromatin assembly factor 1, subunit B (p60)	NM_005441	1.16
dual specificity phosphatase 1	AA530892	1.16
hypothetical protein FLJ22955	NM_024819	1.16
laminin receptor 1 (67kD, ribosomal protein SA)	AL136306	1.16
lysyl oxidase-like 2	BE251211	1.16
Consensus includes gb:AL139349 /DEF=Human DNA sequence from clone RP11-261P9 on chromosome 20. Contains ESTs, STSs, GSSs and a CpG island. Contains the 3 part of the STX16 ge	AL139349	1.16
Ras-related associated with diabetes	NM_004165	1.16
EGF-like-domain, multiple 3	AL134303	1.16
thyrotrophic embryonic factor	U06935	1.16
signal recognition particle receptor ('docking protein')	NM_003139	1.16
nipsnap homolog 1 (C. elegans)	NM_003634	1.16
heme oxygenase (decycling) 2	D21243	1.16
lymphoblastic leukemia derived sequence 1	BC002796	1.16
Consensus includes gb:N36926 /FEA=EST /DB_XREF=gi:1158068 /DB_XREF=est:yy38e04.s1 /CLONE=IMAGE:273534 /UG=Hs.296361 potassium inwardly-rectifying channel,subfamily J, n	N36926	1.16
hypothetical protein	NM_019044	1.16
erythrocyte membrane protein band 4.1-like 3	NM_012307	1.16
sirtuin silent mating type information regulation 2 homolog 6 (S. cerevisiae)	NM_016539	1.16
tripeptidyl peptidase II	AW612376	1.16
KIAA0130 gene product	AI023317	1.16
transformer-2 alpha (htra-2 alpha)	AA831170	1.16
hypothetical protein FLJ20422	NM_017814	1.16
gb:NM_016374.2 /DEF=Homo sapiens RBP1-like protein (BCAA), mRNA. /FEA=mRNA /GEN=BCAA /PROD=RBP1-like protein /DB_XREF=gi:13518237 /FL=gb:NM_016374.2	NM_016374	1.16
glycosylphosphatidylinositol specific phospholipase D1	L11702	1.16
Consensus includes gb:BC005066.1 /DEF=Homo sapiens, clone IMAGE:3347954, mRNA, partial cds. /FEA=mRNA /PROD=Unknown (protein for IMAGE:3347954) /DB_XREF=gi:13477204 /L	BC005066	1.15
solute carrier family 24 (sodium/potassium/calcium exchanger), member 1	AB014602	1.15
diacylglycerol kinase, alpha (80kD)	AF064771	1.15
LDL receptor adaptor protein	AL545035	1.15
myozenin	NM_021245	1.15
Cluster Incl. AI832249:td14g10.x1 Homo sapiens cDNA, 3' end /clone=IMAGE-2075682 /clone_end=3' /gb=AI832249 /gi=5452920 /ug=Hs.211522 /len=545	AI832249	1.15
HLA class II region expressed gene KE2	AW007185	1.15
methyl-CpG binding domain protein 4	AI913365	1.15
hypothetical protein FLJ22582	NM_025045	1.15
similar to KIAA0010 gene product (H. sapiens)	AL096740	1.15
growth differentiation factor 1	NM_001492	1.15
Consensus includes gb:AU147402 /FEA=EST /DB_XREF=gi:11008923 /DB_XREF=est:AU147402 /CLONE=MAMMA1000567 /UG=Hs.287496 Homo sapiens cDNA FLJ12160 fis, clone MAMM	AU147402	1.15
PTD002 protein	NM_016144	1.15
ring finger protein 27	NM_030912	1.15
nuclear receptor subfamily 3, group C, member 1	AI432196	1.15
growth hormone 1	NM_000515	1.15
RAN binding protein 16	NM_015024	1.15
hypothetical protein	AI217472	1.15
inhibitor of Bruton's tyrosine kinase	AF235049	1.15
deformed epidermal autoregulatory factor 1 (Drosophila)	AF068892	1.15
cytochrome P450 retinoid metabolizing protein	NM_019885	1.15
hypothetical protein FLJ10849	AL534972	1.15
apolipoprotein A-I	X02162	1.15
DKFZP564C196 protein	AL050020	1.15
gb:NM_015905.1 /DEF=Homo sapiens transcriptional intermediary factor 1 (TIF1), mRNA. /FEA=mRNA /GEN=TIF1 /PROD=transcriptional intermediary factor 1 alpha /DB_XREF=gi:770623	NM_015905	1.15
Consensus includes gb:AA142929 /FEA=EST /DB_XREF=gi:1712307 /DB_XREF=est:zl40g07.s1 /CLONE=IMAGE:504444 /UG=Hs.288993 ESTs	AA142929	1.15
chromosome 22 open reading frame 3	NM_012265	1.15
hypothetical protein FLJ13491	AI431902	1.15
tripartite motif-containing 2	AK000179	1.15
transposon-derived Buster1 transposase-like protein	NM_021211	1.15
matrin 3	AA129420	1.15
RAB28, member RAS oncogene family	NM_004249	1.15
androgen-induced prostate proliferative shutoff associated protein	NM_015928	1.15
hypothetical protein FLJ10604	NM_018154	1.15
Consensus includes gb:BF448531 /FEA=EST /DB_XREF=gi:11514556 /DB_XREF=est:7n72e02.x1 /CLONE=IMAGE:3570146 /UG=Hs.248979 ESTs, Moderately similar to ALU8_HUMAN ALU	BF448531	1.15

lysosomal-associated protein transmembrane 4 alpha	NM_014713	1.15
H3 histone, family 3B (H3.3B)	BC001124	1.15
paired box gene 8	X69699	1.15
UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase 1 (GalNAc-T1)	AV692127	1.15
7-dehydrocholesterol reductase	NM_001360	1.15
ATP-dependent RNA helicase	BE963238	1.15
cell cycle progression 2 protein	NM_004749	1.15
vascular endothelial growth factor	AF022375	1.15
SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily a, member 3	AI760760	1.15
synaptojanin 2	AF318616	1.15
byrstin-like	NM_004053	1.15
Wiskott-Aldrich syndrome (eczema-thrombocytopenia)	U12707	1.15
heterogeneous nuclear ribonucleoprotein D (AU-rich element RNA binding protein 1, 37kD)	M94630	1.15
v-akt murine thymoma viral oncogene homolog 2	AA769075	1.15
dual-specificity tyrosine-(Y)-phosphorylation regulated kinase 2	NM_006482	1.15
Cip1-interacting zinc finger protein	AF234161	1.15
N-myristoyltransferase 1	AI570834	1.15
Consensus includes gb:AW452623 /FEA=EST /DB_XREF=gi:6993399 /DB_XREF=est:UI-H-BI3-alu-d-09-0-UI.s1 /CLONE=IMAGE:3068608 /UG=Hs.14846 Homo sapiens mRNA; cDNA DKFZp!	AW452623	1.15
KIAA0808 gene product	AI961231	1.15
Janus kinase 3 (a protein tyrosine kinase, leukocyte)	U31601	1.15
homeo box (H6 family) 1	AI939402	1.15
G-protein coupled receptor 88	NM_022049	1.15
like-glycosyltransferase	AB011181	1.15
hypothetical protein	NM_013313	1.15
major histocompatibility complex, class I, F	NM_018950	1.15
singed-like (fascin homolog, sea urchin) (Drosophila)	NM_003088	1.15
protein inhibitor of activated STAT protein PIASy	AK022481	1.15
MCM4 minichromosome maintenance deficient 4 (S. cerevisiae)	AI859865	1.15
paraneoplastic antigen MA1	NM_006029	1.15
postsynaptic protein CRIPT	NM_014171	1.15
hypothetical protein PRO0943	NM_018568	1.15
KIAA1010 protein	AB023227	1.15
hypothetical protein FLJ20344	NM_017776	1.15
KIAA0435 gene product	NM_014801	1.15
hypothetical protein FLJ20113	AL096714	1.15
Consensus includes gb:AA099357 /FEA=EST /DB_XREF=gi:1645257 /DB_XREF=est:zkB5c01.s1 /CLONE=IMAGE:489600 /UG=Hs.15780 ATP-binding cassette, sub-family A (ABC1), membe	AA099357	1.15
hypothetical protein MGC10882	BC004952	1.15
pleckstrin	NM_002664	1.15
steroid sulfatase (microsomal), arylsulfatase C, isozyme S	AU138166	1.15
splicing factor proline/glutamine rich (polypyrimidine tract binding protein associated)	NM_005066	1.15
NIC1-related gene on chromosome 22	BE465318	1.15
KIAA1078 protein	AL110158	1.15
leukocyte membrane antigen	AJ010102	1.15
hypothetical protein FLJ13346	NM_024918	1.15
fumarate hydratase	AI363836	1.15
discs, large (Drosophila) homolog 1	NM_004087	1.15
U5 snRNP-specific 40 kDa protein (hPrp8-binding)	AL157420	1.15
hypothetical protein FLJ20718	NM_017939	1.15
Cluster Incl. AI424872:tg38110.s1 Homo sapiens cDNA, 3' end /clone=IMAGE-2111083 /clone_end=3' /gb=AI424872 /gi=4270895 /ug=Hs.159375 /len=443	AI424872	1.15
sperm associated antigen 11	AF168619	1.15
TBP-interacting protein	NM_018448	1.15
death associated transcription factor 1	NM_022105	1.15
KIAA0332 protein	AB002330	1.15
DKFZP564K247 protein	NM_014056	1.15
5-hydroxytryptamine (serotonin) receptor 5A	NM_024012	1.15
eukaryotic translation initiation factor 4B	NM_001417	1.15
hypothetical protein FLJ11132	NM_018335	1.15
huntingtin interacting protein 1	NM_005338	1.15
ATPase, H+ transporting, lysosomal (vacuolar proton pump) 9kD	NM_003945	1.15
ATPase, Ca++ transporting, plasma membrane 1	M95541	1.15
sudD (suppressor of bimD6, Aspergillus nidulans) homolog	AW006290	1.15
gb:NM_016104.1 /DEF=Homo sapiens PTD013 protein (PTD013), mRNA. /FEA=mRNA /GEN=PTD013 /PROD=PTD013 protein /DB_XREF=gi:7706668 /UG=Hs.279857 PTD013 protein /FL	NM_016104	1.15
solute carrier family 18 (vesicular monoamine), member 2	AI269290	1.15

Consensus includes gb:R27378 /FEA=EST /DB_XREF=gi:783513 /DB_XREF=est:yh53g11.r1 /CLONE=IMAGE:133508 /UG=Hs.322426 ESTs, Moderately similar to TIAR_HUMAN NUCLEOLYSI	R27378	1.15
CGI-105 protein	AA723370	1.14
ganglioside-induced differentiation-associated protein 1	NM_018972	1.14
GM2 ganglioside activator protein	X61094	1.14
BAI1-associated protein 2	AB017120	1.14
mitogen-activated protein kinase 3	X60188	1.14
A kinase (PRKA) anchor protein 10	AA456929	1.14
hypothetical protein 628	BC000974	1.14
transmembrane 4 superfamily member 6	U84895	1.14
nuclear receptor coactivator 4	AL162047	1.14
plakophilin 3	AF053719	1.14
mannose-6-phosphate receptor (cation dependent)	NM_002355	1.14
GW128 protein	BF246499	1.14
sialidase 2 (cytosolic sialidase)	NM_005383	1.14
polymerase (DNA-directed), alpha (70kD)	NM_002689	1.14
transcription factor 3 (E2A immunoglobulin enhancer binding factors E12/E47)	X52078	1.14
Consensus includes gb:AA912711 /FEA=EST /DB_XREF=gi:3052103 /DB_XREF=est:ol30f08.s1 /CLONE=IMAGE:1524999 /UG=Hs.26395 erythrocyte membrane protein band 4.1-like 1	AA912711	1.14
CDP-diacylglycerol synthase (phosphatidate cytidylyltransferase) 2	AL568982	1.14
gb:L08666.1 /DEF=Homo sapiens porin (por) mRNA, complete cds and truncated cds. /FEA=mRNA /GEN=por; por /PROD=porin; porin /DB_XREF=gi:190199 /FL=gb:L08666.1	L08666	1.14
Consensus includes gb:BE782754 /FEA=EST /DB_XREF=gi:10203952 /DB_XREF=est:601471903F1 /CLONE=IMAGE:3874921 /UG=Hs.102178 syntaxin 16 /FL=gb:AF008936.1	BE782754	1.14
histone acetyltransferase	NM_007067	1.14
KIAA1034 protein	AB028957	1.14
DKFZP4348168 protein	AL133049	1.14
reproduction 8	D83768	1.14
similar to S. cerevisiae SSM4	AK000970	1.14
SEC24 related gene family, member A (S. cerevisiae)	BE645231	1.14
growth hormone 1	NM_022560	1.14
dCMP deaminase	NM_001921	1.14
MIK-related kinase epsilon; inducible IkappaB kinase	NM_014002	1.14
hypothetical protein	AK025647	1.14
DKFZP434N043 protein	BG538800	1.14
hypothetical protein FLJ20450	NM_017827	1.14
ubiquitin specific protease 15	AF153604	1.14
protease, serine, 1 (trypsin 1)	U66061	1.14
SH3-domain GRB2-like 2	NM_003026	1.14
SELENOPHOSPHATE SYNTHETASE ; Human selenium donor protein	BC000941	1.14
leukocyte immunoglobulin-like receptor, subfamily A (with TM domain), member 2	U82276	1.14
aldo-keto reductase family 1, member C3 (3-alpha hydroxysteroid dehydrogenase, type II)	AB018580	1.14
butyrophilin, subfamily 2, member A1	U90142	1.14
ATP synthase, H+ transporting, mitochondrial F0 complex, subunit c (subunit 9) isoform 3	NM_001689	1.14
chromosome 1 open reading frame 16	BF509099	1.14
F-box only protein 5	NM_012177	1.14
Consensus includes gb:AL110206.1 /DEF=Homo sapiens mRNA; cDNA DKFZp586N2022 (from clone DKFZp586N2022). /FEA=mRNA /DB_XREF=gi:5817125 /UG=Hs.306339 Homo sapiens	AL110206	1.14
KIAA1453 protein	NM_025090	1.14
iroquois homeobox protein 5	U90304	1.14
putative DNA binding protein	NM_007358	1.14
hypothetical protein FLJ11370	NM_024961	1.14
hypothetical protein FLJ13848	NM_024771	1.14
uncharacterized hypothalamus protein HT012	NM_018473	1.14
hypothetical protein FLJ20174	NM_017699	1.14
ribulose-5-phosphate-3-epimerase	BE964473	1.14
lysophospholipase II	NM_007260	1.14
hypothetical protein FLJ20641	NM_017915	1.14
transport-secretion protein 2.2,	AF055000	1.14
tumor necrosis factor, alpha-induced protein 3	NM_006290	1.14
ATP-binding cassette, sub-family B (MDR/TAP), member 1	AF016535	1.14
zinc finger protein 313	NM_018683	1.14
ring finger protein 34	NM_025126	1.14
golgi transport complex 1 (90 kDa subunit)	AU152134	1.14
Homer, neuronal immediate early gene, 3	AC002985	1.14
KIAA0229 protein	AI990523	1.14
related RAS viral (r-ras) oncogene homolog 2	AI431643	1.14
Rh type C glycoprotein	NM_016321	1.14

protein kinase C, zeta	NM_002744	1.14
uncharacterized hematopoietic stem/progenitor cells protein MDS030	NM_018465	1.14
hypothetical protein FLJ10483	NM_018108	1.14
hypothetical protein FLJ20596	NM_017895	1.14
sarcoglycan, epsilon	NM_003919	1.14
hypothetical protein FLJ22724	NM_024532	1.14
KIAA1089 protein	AB029012	1.14
neural proliferation, differentiation and control, 1	NM_015392	1.14
Cluster Incl. AW026481:ew14a04.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-990913 /clone_end=3 /gb=AW026481 /gi=5880011 /ug=Hs.48396 /len=496	AW026481	1.14
nucleosome assembly protein 1-like 1	AW148801	1.14
hypothetical protein	AW873594	1.14
UDP glycosyltransferase 1 family, polypeptide A6	NM_001072	1.14
TTK protein kinase	NM_003318	1.14
SHP2 interacting transmembrane adaptor	NM_014450	1.14
spectrin SH3 domain binding protein 1	AF006516	1.14
dual specificity phosphatase 6	BC005047	1.14
platelet-derived growth factor receptor-like	NM_006207	1.14
glioma tumor suppressor candidate region gene 1	NM_015711	1.14
hypothetical protein FLJ10688	NM_018179	1.14
KIAA1922 protein	AK026910	1.14
xylosylprotein beta1,4-galactosyltransferase, polypeptide 7 (galactosyltransferase I)	AK022566	1.14
myosin, heavy polypeptide 11, smooth muscle	S67238	1.14
solute carrier family 7 (cationic amino acid transporter, y+ system), member 7	NM_003982	1.14
hypothetical protein PRO0758	NM_018563	1.14
Consensus includes gb:AW303476 /FEA=EST /DB_XREF=gi:6713165 /DB_XREF=est:zv19e10.x1 /CLONE=IMAGE:2813610 /UG=Hs.132299 ESTs	AW303476	1.14
heterogeneous nuclear ribonucleoprotein U (scaffold attachment factor A)	NM_004501	1.14
RAD52 homolog (S. cerevisiae)	NM_002879	1.14
Cluster Incl. AI144299:qb59h06.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-1704443 /clone_end=3 /gb=AI144299 /gi=3666108 /ug=Hs.106843 /len=765	AI144299	1.14
HMBA-inducible	AW193511	1.14
Consensus includes gb:BE299495 /FEA=EST /DB_XREF=gi:9183243 /DB_XREF=est:600944774T1 /CLONE=IMAGE:2960610 /UG=Hs.218329 hypothetical protein /FL=gb:NM_015383.1	BE299495	1.14
mitogen-activated protein kinase 4	BF115223	1.14
PCTAIRE protein kinase 1	BC001048	1.14
phosducin-like	AF031463	1.14
Consensus includes gb:AU148040 /FEA=EST /DB_XREF=gi:11009561 /DB_XREF=est:AU148040 /CLONE=MAMMA1002428 /UG=Hs.301615 Homo sapiens cDNA FLJ12370 fis, clone MAMM	AU148040	1.14
UDP glycosyltransferase 8 (UDP-galactose ceramide galactosyltransferase)	NM_003360	1.14
ribokinase	NM_022128	1.14
hypothetical protein FLJ10525	NM_018126	1.14
hypothetical protein FLJ22613	NM_024777	1.14
vaccinia related kinase 2	NM_006296	1.14
glycine cleavage system protein H (aminomethyl carrier)	AW237404	1.14
hypothetical protein FLJ20539	NM_017870	1.14
activating transcription factor 5	BC005174	1.14
nucleobindin 1	NM_006184	1.14
LIM domain-containing preferred translocation partner in lipoma	BF221852	1.14
fucosyltransferase 2 (secretor status included)	NM_000511	1.14
activation-induced cytidine deaminase	NM_020661	1.14
TNF receptor-associated factor 3	AI721219	1.14
xeroderma pigmentosum, complementation group A	NM_000380	1.14
Consensus includes gb:AL036840 /FEA=EST /DB_XREF=gi:5406335 /DB_XREF=est:DKFZp564N1963_s1 /CLONE=DKFZp564N1963 /UG=Hs.22370 Homo sapiens mRNA; cDNA DKFZp5640	AL036840	1.14
phosphatidylinositol glycan, class B	NM_004855	1.14
hypothetical protein MGC14817	AF255792	1.13
protein kinase C binding protein 1	BC001004	1.13
complement component 1, q subcomponent, beta polypeptide	NM_000491	1.13
transferrin receptor 2	NM_003227	1.13
interleukin 12 receptor, beta 1	NM_005535	1.13
phosphatidylcholine transfer protein	NM_021213	1.13
laminin, beta 3 (nicein (125kD), kalinin (140kD), BM600 (125kD))	L25541	1.13
fascin (Strongylocentrotus purpuratus) homolog 2 (actin-bundling protein, retinal)	NM_012418	1.13
CDC10 cell division cycle 10 homolog (S. cerevisiae)	AU157515	1.13
complement component 4B	AL049547	1.13
diphtheria toxin receptor (heparin-binding epidermal growth factor-like growth factor)	BE222436	1.13
KIAA1055 protein	BF195608	1.13
hypothetical protein MGC955	W37846	1.13

claudin 11 (oligodendrocyte transmembrane protein)	NM_005602	1.13
mitogen-activating protein kinase kinase kinase kinase 2	NM_004579	1.13
glutaminyl-peptide cyclotransferase (glutaminyl cyclase)	NM_012413	1.13
6 protein-coupled receptor 8	NM_005286	1.13
eukaryotic translation initiation factor SA2	NM_020390	1.13
progastagen-associated endometrial protein (placental protein 14, pregnancy-associated endometrial alpha-2-globulin, alpha uterine protein)	NM_002571	1.13
nucleolar protein ANKT	NM_016359	1.13
26S proteasome-associated pad1 homolog	NM_005805	1.13
solute carrier family 29 (nucleoside transporters), member 2	AF034102	1.13
hypothetical protein FLJ11342	NM_018394	1.13
DKFZP434F195 protein	AL043774	1.13
UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase 7 (GalNAc-T7)	NM_017423	1.13
SUMO-1 activating enzyme subunit 2	NM_005499	1.13
active BCR-related gene	AL527773	1.13
upstream transcription factor 2, c-fos interacting	AY007087	1.13
Consensus includes gb:AF007132.1 /DEF=Homo sapiens clone 23551 mRNA sequence. /FEA=mRNA /DB_XREF=gi:2852606 /UG=Hs.184019 Homo sapiens clone 23551 mRNA sequence	AF007132	1.13
transforming, acidic coiled-coil containing protein 1	NM_006283	1.13
hypothetical protein FLJ20730	NM_017945	1.13
keratin, hair, acidic,38	NM_002279	1.13
peroxisomal biogenesis factor 16	BC000467	1.13
T cell receptor beta locus	AF009660	1.13
forkhead box O38	AF032887	1.13
interleukin 1, alpha	M15329	1.13
POU domain, class 5, transcription factor 1	AF268617	1.13
hypothetical protein FLJ14154	AI246641	1.13
KIAA0592 protein	AV728658	1.13
titin	NM_003319	1.13
KIAA0376 protein	AB002374	1.13
hypothetical protein FLJ20315	NM_017763	1.13
solute carrier family 35 (UDP-N-acetylglucosamine (UDP-GlcNAc) transporter), member 3	BC005136	1.13
ras-related C3 botulinum toxin substrate 2 (rho family, small GTP binding protein Rac2)	NM_002872	1.13
actin	AA745954	1.13
hypothetical protein FLJ20279	NM_017740	1.13
tektin 2 (testicular)	AB033823	1.13
ATPase, Na+/K+ transporting, alpha 3 polypeptide	NM_000703	1.13
HIV-1 rev binding protein 2	AI950314	1.13
cytokine-like nuclear factor n-pac	BC003693	1.13
HIV TAT specific factor 1	NM_014500	1.13
tetraspan 3	BC000704	1.13
synaptotagmin V	AI659957	1.13
kinesin-like 3	NM_005355	1.13
hypothetical protein MGC2479	NM_024299	1.13
hypothetical C2H2 zinc finger protein FLJ22504	NM_022095	1.13
ADP-ribosylation factor 1	AF052179	1.13
enhancer of zeste homolog 1 (Drosophila)	AB002386	1.13
Consensus includes gb:AK025191.1 /DEF=Homo sapiens cDNA: FLJ21538 fis, clone COL06151. /FEA=mRNA /DB_XREF=gi:10437654 /UG=Hs.306783 Homo sapiens cDNA: FLJ21538 fis, cl	AK025191	1.13
centaurin, alpha 1	AW050627	1.13
aconitase 1, soluble	NM_002197	1.13
heat shock transcription factor 2	BC005329	1.13
ribosomal protein S6 kinase, 90kD, polypeptide 2	BC002363	1.13
DKFZP566C243 protein	AK021433	1.13
jun D proto-oncogene	AI339541	1.13
hypothetical protein FLJ22439	AW027923	1.13
phosphatidylinositol glycan, class H	BC004100	1.13
small nuclear ribonucleoprotein polypeptide B"	NM_003092	1.13
retinoid x receptor interacting protein	NM_016290	1.13
keratin 8	U76549	1.13
phospholipase C, beta 2	NM_004573	1.13
early development regulator 2 (homolog of polyhomeotic 2)	NM_004427	1.13
Consensus includes gb:BG031677 /FEA=EST /DB_XREF=gi:12422203 /DB_XREF=est:602300830F1 /CLONE=IMAGE:4402326 /UG=Hs.58210 ESTs	BG031677	1.13
gb:AB058895.1 /DEF=Homo sapiens GHRHRpsv mRNA for growth hormone releasing hormone receptor pituitary splice variant, complete cds. /FEA=mRNA /GEN=GHRHRpsv /PROD=growth h	AB058895	1.13
ubiquitin-conjugating enzyme E2L 3	NM_003347	1.13
metazin 2	NM_006554	1.13

hypothetical protein FLJ23471	NM_024723	1.13
fibroblast growth factor 22	NM_020637	1.13
uroporphyrinogen decarboxylase	M14016	1.13
SUMO-1-specific protease	NM_015571	1.13
ATP-binding cassette, sub-family C (CFTR/MRP), member 1	NM_004996	1.13
Consensus includes gb:AA543076 /FEA=EST /DB_XREF=gi:2291556 /DB_XREF=est:nf96f02.s1 /CLONE=IMAGE:927771 /UG=Hs.82171 Homo sapiens (clone 115392) mRNA	AA543076	1.13
mannosidase, alpha, class 1B, member 1	NM_016219	1.13
protein tyrosine phosphatase-like (proline instead of catalytic arginine), member a	NM_014241	1.13
SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily c, member 1	AL040633	1.13
testis expressed sequence 138	NM_031273	1.13
hypothetical protein BC002942	AI081535	1.13
hypothetical protein FLJ14360	N38751	1.13
interleukin 19	NM_013371	1.13
hypothetical protein FLJ20354	NM_017779	1.13
homolog of mouse quaking QKI (KH domain RNA binding protein)	AI114716	1.13
hypothetical protein MGC2776	NM_025265	1.13
c-myc promoter-binding protein	BE268538	1.13
HBV associated factor	NM_006462	1.13
homolog of yeast MOG1	NM_014185	1.13
chondroitin sulfate proteoglycan 4 (melanoma-associated)	BE857703	1.13
Deleted in split-hand/split-foot 1 region	NM_006304	1.13
dysferlin, limb girdle muscular dystrophy 2B (autosomal recessive)	NM_003494	1.13
hypothetical protein FLJ11786	NM_030892	1.13
hypothetical protein STRAT11499	NM_021242	1.13
ATPase, (Na+)/K+ transporting, beta 4 polypeptide	NM_012069	1.13
junctional adhesion molecule 1	AF154005	1.13
Consensus includes gb:AK023938.1 /DEF=Homo sapiens cDNA FLJ13876 fis, clone THYRO1001401. /FEA=mRNA /DB_XREF=gi:10436033 /UG=Hs.287604 Homo sapiens cDNA FLJ13876 fi	AK023938	1.13
T-cell, immune regulator 1	NM_006019	1.13
Consensus includes gb:BG390664 /FEA=EST /DB_XREF=gi:13284112 /DB_XREF=est:602415258F1 /CLONE=IMAGE:4523487 /UG=Hs.97496 YY1 transcription factor /FL=gb:M77698.1 gb	BG390664	1.13
Consensus includes gb:AF056433.1 /DEF=Homo sapiens clone FBD3 Cri-du-chat critical region mRNA. /FEA=mRNA /DB_XREF=gi:3044151 /UG=Hs.8078 Homo sapiens clone FBD3 Cri-du-c	AF056433	1.13
enhancer of filamentation 1 (cas-like docking; Crk-associated substrate related)	U64317	1.13
fibulin 1	NM_006486	1.13
hypothetical protein FLJ11040	NM_018307	1.13
nasopharyngeal epithelium specific protein 1	NM_012337	1.13
disrupter of silencing 10	BC004546	1.13
PR domain containing 4	W22625	1.13
PTD015 protein	NM_014040	1.13
syntrophin, beta 2 (dystrophin-associated protein A1, 59kD, basic component 2)	AA741303	1.13
SWAP-70 protein	BC000134	1.13
haptoglobin-related protein	NM_020995	1.13
endoplasmic reticulum chaperone SIL1, homolog of yeast	NM_022464	1.13
Consensus includes gb:AL008730 /DEF=Human DNA sequence from PAC 48717 on chromosome 6q21-22.1. Contains an unknown gene coding for three alternative mRNAs. Contains ESTs, ST	AL008730	1.13
HSPC047 protein	NM_014147	1.13
fumarylacetoacetate hydrolase (fumarylacetoacetase)	NM_000137	1.13
peroxisomal biogenesis factor 16	AB016531	1.13
potassium voltage-gated channel, KQT-like subfamily, member 2	NM_004518	1.13
Consensus includes gb:Z98950 /DEF=Human DNA sequence from PAC 507115 on chromosome Xq26.3-27.3. Contains 60S ribosomal protein L44 (L41, L36) like gene, ESTs, STSs and a poly	Z98950	1.13
SUMO-1-specific protease	AK001406	1.13
craniofacial development protein 1	NM_006324	1.13
chromosome 5 open reading frame 4	H93077	1.13
gb:BC002877.1 /DEF=Homo sapiens, Similar to hypothetical protein FLJ11585, clone MGC:11258, mRNA, complete cds. /FEA=mRNA /PROD=Similar to hypothetical protein FLJ11585 /DB_X	BC002877	1.13
insulin-like growth factor 2 receptor	NM_000876	1.13
KIAA1069 protein	AK022610	1.13
FK506 binding protein 1B (12.6 kD)	AF322070	1.13
hypothetical protein FLJ20069	NM_017651	1.13
interferon, alpha 17	M38289	1.13
neurexin 2	AB035266	1.12
cysteine-rich protein 2	U36190	1.12
TATA box binding protein	NM_003194	1.12
methyl-Cp6 binding domain protein 4	AI913365	1.12
hypothetical protein FLJ10582	NM_018147	1.12
hypothetical protein FLJ12085	NM_022771	1.12
splicing factor (CC1.3)	AI890947	1.12

phosphoribosyl pyrophosphate synthetase-associated protein 1	NM_002766	1.12
KIAA0993 protein	AL536319	1.12
RAB2, member RAS oncogene family	AI189609	1.12
low density lipoprotein receptor-related protein 8, apolipoprotein e receptor	NM_004631	1.12
diaphorase (NADH) (cytochrome b-5 reductase)	NM_000398	1.12
lanosterol synthase (2,3-oxidosqualene-lanosterol cyclase)	D63807	1.12
KIAA0931 protein	AB023148	1.12
KIAA1039 protein	AK000478	1.12
YY1 transcription factor	AA748649	1.12
protein phosphatase 2 (formerly 2A), regulatory subunit B" (PR 72), alpha isoform and (PR 130), beta isoform	NM_002718	1.12
integral inner nuclear membrane protein	NM_014319	1.12
hypothetical protein FLJ22329	NM_024656	1.12
Fanconi anemia, complementation group A	NM_000135	1.12
suppressor of Ty 4 homolog 1 (S. cerevisiae)	NM_003168	1.12
IQ motif containing GTPase activating protein 1	NM_003870	1.12
Consensus includes gb:AL354872 /DEF=Human DNA sequence from clone RP11-42015 on chromosome 1. Contains ESTs, STSs, GSSs and a CpG island. Contains the CTH gene for two isofon	AL354872	1.12
DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 15	NM_001358	1.12
SEC7 homolog	AK023421	1.12
protein phosphatase methylesterase-1	AL040538	1.12
lysozyme homolog	NM_020426	1.12
hypothetical protein FLJ20036	NM_017632	1.12
phosphodiesterase 8A	W73272	1.12
crystallin, zeta (quinone reductase)	NM_001889	1.12
thymidylate synthetase	NM_001071	1.12
KIAA1068 protein	NM_015332	1.12
solute carrier family 25 (mitochondrial carrier, brain), member 14	AF155810	1.12
KIAA0179 protein	AI400490	1.12
CGI-136 protein	NM_016069	1.12
homeo box B7	NM_004502	1.12
DKFZP434F195 protein	AA019977	1.12
hypothetical protein FLJ10008	NM_017970	1.12
vitiligo-associated protein VIT-1	AL117620	1.12
SAR1 protein	AY008268	1.12
COX11 homolog, cytochrome c oxidase assembly protein (yeast)	BC005895	1.12
actin cross-linking factor	W61052	1.12
arginine vasopressin receptor 2 (nephrogenic diabetes insipidus)	AF030626	1.12
hypothetical protein FLJ22169	NM_024085	1.12
cytochrome b5 reductase 1 (B5R.1)	NM_016243	1.12
iduronate 2-sulfatase (Hunter syndrome)	AF050145	1.12
Consensus includes gb:AL133015.1 /DEF=Homo sapiens mRNA; cDNA DKFZp43402417 (from clone DKFZp43402417); partial cds. /FEA=mRNA /GEN=DKFZp43402417 /PROD=hypothetic	AL133015	1.12
Consensus includes gb:AI983043 /FEA=EST /DB_XREF=gi:5810262 /DB_XREF=est:wz30b11.x1 /CLONE=IMAGE:2559549 /UG=Hs.246885 hypothetical protein FLJ20783 /FL=gb:NM_017!	AI983043	1.12
lysosomal-associated membrane protein 2	J04183	1.12
RAS p21 protein activator 2	NM_006506	1.12
CAP-binding protein complex interacting protein 2	BF939788	1.12
Cluster Incl. AI302185:qn58d02.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-1902435 /clone_end=3 /gb=AI302185 /gi=3961531 /ug=Hs.4789 /len=522	AI302185	1.12
KIAA0397 gene product	AB007857	1.12
COP9 (constitutive photomorphogenic, Arabidopsis, homolog) subunit 3	NM_003653	1.12
CGI-63 protein	NM_016011	1.12
annexin A2	BE908217	1.12
Golgi apparatus protein 1	NM_012201	1.12
autocrine motility factor receptor	AF124145	1.12
ring finger protein 14	AB022663	1.12
E1B-55kDa-associated protein 5	BC004242	1.12
hypothetical gene BC008967	BE299456	1.12
Consensus includes gb:U50277.1 /DEF=Human breast cancer suppressor element Ishmael Upper CP1 mRNA, partial cds. /FEA=mRNA /PROD=suppressor element Ishmael Upper CP1 /DB_X	U50277	1.12
diazepam binding inhibitor (GABA receptor modulator, acyl-Coenzyme A binding protein)	NM_020548	1.12
hypothetical protein FLJ10715	NM_018190	1.12
exosome component Rrp41	AI571298	1.12
anti-oxidant protein 2 (non-selenium glutathione peroxidase, acidic calcium-independent phospholipase A2)	NM_004905	1.12
protein phosphatase 1, regulatory (inhibitor) subunit 12A	NM_002480	1.12
KIAA1696 protein	NM_016621	1.12
glycophorin C (Gerbich blood group)	NM_002101	1.12
Snf2-related CBP activator protein	AB002307	1.12

DKFZP564C152 protein	AL049980	1.12
tripartite motif-containing 14	NM_014788	1.12
TRIAD3 protein	BC000787	1.12
karyopherin (importin) beta 2	AI653355	1.12
E74-like factor 1 (ets domain transcription factor)	AL559590	1.12
phosphoprotein enriched in astrocytes 15	NM_003768	1.12
double homeobox, 2	NM_012147	1.12
forkhead box O3A	NM_001455	1.12
nucleolar cysteine-rich protein	H82458	1.12
sarcoglycan, delta (35kD dystrophin-associated glycoprotein)	U58331	1.12
DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 28	NM_018380	1.12
melan-A	NM_005511	1.12
solute carrier family 2, (facilitated glucose transporter) member 8	NM_014580	1.12
Consensus includes gb:AL078636.1 /DEF=Homo sapiens mRNA full length insert cDNA clone EUROIMAGE 117929. /FEA=mRNA /DB_XREF=gi:5051866 /UG=Hs.227171 Homo sapiens mRN	AL078636	1.12
KIAA0268 protein	D87742	1.12
adaptor-related protein complex 3, mu 2 subunit	NM_006803	1.12
HLA-B associated transcript 2	NM_004638	1.12
hypothetical protein FLJ20511	NM_017853	1.12
KIAA0680 gene product	NM_014721	1.12
UDP-glucuronic acid/UDP-N-acetylgalactosamine dual transporter	N80922	1.12
airway trypsin-like protease	NM_004262	1.12
H2B histone family, member B	BC002842	1.12
KIAA0202 protein	BC001329	1.12
similar to KIAA0010 gene product (H. sapiens)	BE856776	1.12
zinc finger protein 174	BC001161	1.12
KIAA0015 gene product	D13640	1.12
phosphotidylinositol transfer protein	H15647	1.12
acyl-Coenzyme A oxidase 2, branched chain	NM_003500	1.12
cyclin G1	BC000196	1.12
hypothetical protein FLJ10379	NM_018079	1.12
hypothetical protein DKFZp434I1916	AL136771	1.12
Wolf-Hirschhorn syndrome candidate 2	AF101434	1.12
lysophospholipase II	AK024724	1.12
chromosome 11 open reading frame 17	NM_020642	1.12
adducin 3 (gamma)	AI818488	1.12
sigma receptor (SR31747 binding protein 1)	AF226604	1.12
phosphatidylinositol-4-phosphate 5-kinase, type I, beta	U78581	1.12
retinoblastoma 1 (including osteosarcoma)	NM_000321	1.12
NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 2 (8kD, AGGG)	NM_004546	1.12
Sec23 homolog B (S. cerevisiae)	BC005032	1.12
proteasome (prosome, macropain) 26S subunit, non-ATPase, 4	NM_002810	1.12
KIAA0828 protein	AB020635	1.12
NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 6 (17kD, B17)	NM_002493	1.12
GTPase regulator associated with the focal adhesion kinase pp125	BE671084	1.12
hypothetical protein DT1P1A10	AA223871	1.12
splicing factor, arginine/serine-rich 9	NM_003769	1.12
intercellular adhesion molecule 1 (CD54), human rhinovirus receptor	AI608725	1.12
solute carrier family 5 (sodium/glucose cotransporter), member 2	NM_003041	1.12
KIAA0923 protein	AK001710	1.12
sphingosine kinase type 2 isoform	AF245447	1.12
FtsJ homolog 1 (E. coli)	NM_012280	1.12
CD2-associated protein	NM_012120	1.12
SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily c, member 1	AA593983	1.12
isoleucine-tRNA synthetase	NM_013417	1.12
golgi autoantigen, golgin subfamily b, macrogolgin (with transmembrane signal), 1	NM_004487	1.12
lymphoid blast crisis oncogene	NM_006738	1.12
lymphocyte activation-associated protein	NM_015990	1.12
biliverdin reductase A	AA740186	1.12
phosphoserine aminotransferase	W72527	1.12
KIAA0970 protein	NM_014923	1.12
KIAA0788 protein	BC001417	1.12
RING1 and YY1 binding protein	AB029551	1.12
breakpoint cluster region	NM_004327	1.12

microfibrillar-associated protein 4	R72286	1.12
RAB3B, member RAS oncogene family	NM_002867	1.12
ligase III, DNA, ATP-dependent	NM_013975	1.12
hypothetical protein	NM_020313	1.12
small nuclear RNA activating complex, polypeptide 5, 19kD	BE259395	1.12
24 kDa intrinsic membrane protein	BF057649	1.12
Consensus includes gb:AK026188.1 /DEF=Homo sapiens cDNA: FLJ22535 fis, clone HRC13115, highly similar to AF152336 Homo sapiens protocadherin gamma B7 (PCDH-gamma-B7) mRNA	AK026188	1.12
kinetochore associated 1	NM_014708	1.12
syntrophin, beta 1 (dystrophin-associated protein A1, 59kD, basic component 1)	BG484314	1.12
KIAA0317 gene product	NM_014821	1.12
disrupted in bipolar disorder 1	NM_024740	1.12
CTD (carboxy-terminal domain, RNA polymerase II, polypeptide A) phosphatase, subunit 1	NM_004715	1.12
immunoglobulin (CD79A) binding protein 1	NM_001551	1.12
cullin 4A	NM_003589	1.12
leukotriene A4 hydrolase	J02959	1.12
mannan-binding lectin serine protease 2	AB008047	1.12
dihydrolipoamide S-acetyltransferase (E2 component of pyruvate dehydrogenase complex)	AW299740	1.12
glutamate receptor, ionotropic, N-methyl D-aspartate 1	L05666	1.12
glucosidase, beta; acid (includes glucosylceramidase)	K02920	1.11
hypothetical protein DKFZp434G1415	NM_031292	1.11
putative UDP-GalNAc:polypeptide N-acetylgalactosaminyltransferase T9	NM_017540	1.11
ATP-binding cassette, sub-family G (WHITE), member 5 (sterolin 1)	NM_022436	1.11
KIAA0329 gene product	NM_014844	1.11
protease, serine, 22	NM_006025	1.11
candidate mediator of the p53-dependent G2 arrest	NM_019845	1.11
hypothetical protein FLJ21816	NM_024675	1.11
Consensus includes gb:AK023081.1 /DEF=Homo sapiens cDNA FLJ13019 fis, clone NT2RP3000736, highly similar to Human mRNA for KIAA0140 gene. /FEA=mRNA /DB_XREF=gi:1043483	AK023081	1.11
solute carrier family 29 (nucleoside transporters), member 1	NM_004955	1.11
hypothetical protein PRO2015	NM_018512	1.11
protein phosphatase 1, regulatory (inhibitor) subunit 16B	AB020630	1.11
Consensus includes gb:AK024135.1 /DEF=Homo sapiens cDNA FLJ14073 fis, clone HEMBB1001812. /FEA=mRNA /DB_XREF=gi:10436441 /UG=Hs.296751 Homo sapiens cDNA FLJ14073 f	AK024135	1.11
hypothetical protein FLJ12584	NM_025139	1.11
Consensus includes gb:AA477655 /FEA=EST /DB_XREF=gi:2206289 /DB_XREF=est:zu37h09.s1 /CLONE=IMAGE:740225 /UG=Hs.181307 H3 histone, family 3A	AA477655	1.11
epimorphin	NM_001980	1.11
5'-nucleotidase (purine), cytosolic type B	BC001595	1.11
putative secreted protein	NM_015926	1.11
Down syndrome critical region gene 1-like 2	NM_013441	1.11
synaptosomal-associated protein, 23kD	U55936	1.11
golgi autoantigen, golgin subfamily a, 3	NM_005895	1.11
olfactomedin 1	NM_006334	1.11
TNFRSF1A-associated via death domain	N36774	1.11
tumor necrosis factor receptor superfamily, member 6b, decoy	AL121845	1.11
xeroderma pigmentosum, complementation group C	D21089	1.11
HSPC034 protein	AL031427	1.11
hypothetical protein FLJ23027	AA524500	1.11
hypothetical protein FLJ10392	NM_018084	1.11
nuclear pore complex protein	NM_020401	1.11
proliferation-associated 2G4, 38kD	U87954	1.11
zinc finger protein 189	NM_003452	1.11
Bicaudal D homolog 1 (Drosophila)	NM_001714	1.11
neuron-specific protein	BC001745	1.11
nucleoporin 88kD	NM_002532	1.11
hypothetical protein FLJ13956	NM_024707	1.11
methylmalonyl Coenzyme A mutase	AI433712	1.11
collagen, type VI, alpha 1	AI141603	1.11
angiotensin I converting enzyme (peptidyl-dipeptidase A) 1	AI623989	1.11
astrotactin 2	AK024064	1.11
FtsJ homolog 3 (E. coli)	NM_017647	1.11
hypothetical protein MGC11138	AI675178	1.11
hypothetical protein DC50	NM_031210	1.11
homeo box B5	AI052747	1.11
Kruppel-like factor 12	NM_016285	1.11
phospholipase A2, group IVB (cytosolic)	NM_005090	1.11

glutamate rich WD repeat protein GRWD	AF337808	1.11
choline kinase-like	NM_005198	1.11
hypothetical protein FLJ10902	NM_018266	1.11
KIAA0461 protein	AB007930	1.11
chromosome 22 open reading frame 4	BC001292	1.11
potassium inwardly-rectifying channel, subfamily J, inhibitor 1	NM_002244	1.11
putative zinc finger protein from EUROIMAGE 566589	AL121585	1.11
KIAA1085 protein	R39094	1.11
zinc finger protein, X-linked	NM_003410	1.11
hypothetical protein FLJ12949	NM_023008	1.11
sarcoglycan, beta (43kD dystrophin-associated glycoprotein)	U29586	1.11
PTPRF interacting protein, binding protein 2 (liprin beta 2)	AI692180	1.11
testis-specific transcript, Y-linked 1	AF000990	1.11
IM4 protein	NM_007213	1.11
DKFZP58680923 protein	AB033105	1.11
activated leucocyte cell adhesion molecule	BF242905	1.11
KIAA0440 protein	NM_015556	1.11
collagen, type VI, alpha 2	AY029208	1.11
centaurin, delta 1	AK023737	1.11
reticulon 3	NM_006054	1.11
KIAA1232 protein	AI916274	1.11
vesicle-associated membrane protein 2 (synaptobrevin 2)	AI955119	1.11
Consensus includes gb:AI693140 /FEA=EST /DB_XREF=gi:4970480 /DB_XREF=est:wd68a05.x1 /CLONE=IMAGE:2336720 /UG=Hs.22907 Homo sapiens mRNA; cDNA DKFZp586F071 (from	AI693140	1.11
Janus kinase 1 (a protein tyrosine kinase)	AL039831	1.11
heterogeneous nuclear ribonucleoprotein D (AU-rich element RNA binding protein 1, 37kD)	W74620	1.11
hypothetical protein FLJ21802	NM_024644	1.11
beta-2-microglobulin	NM_004048	1.11
protein phosphatase	NM_016364	1.11
DKFZP586F1918 protein	AL050091	1.11
glycosyltransferase AD-017	NM_018446	1.11
reversion-inducing-cysteine-rich protein with kazal motifs	AK022897	1.11
ATP synthase, H+ transporting, mitochondrial F1 complex, gamma polypeptide 1	BC000931	1.11
hypothetical protein similar to ankyrin repeat-containing priotein AKR1	NM_019028	1.11
CLIP-associating protein 2	AV725315	1.11
myosin 5C	NM_018728	1.11
insulin-like growth factor 2 receptor	BG031974	1.11
LIM protein (similar to rat protein kinase C-binding enigma)	AA196325	1.11
solute carrier family 25 (mitochondrial carrier, Aralar), member 12	AI887457	1.11
v-jun sarcoma virus 17 oncogene homolog (avian)	BG491844	1.11
aryl hydrocarbon receptor nuclear translocator	AF001307	1.11
exonuclease NEF-sp	NM_030941	1.11
hypothetical protein FLJ23053	NM_022907	1.11
hypothetical protein FLJ14360	BF732879	1.11
calcitonin-related polypeptide, beta	AA747379	1.11
hypothetical protein	NM_014306	1.11
CGI-01 protein	NM_015935	1.11
PHD finger protein 1	AL050332	1.11
lung cancer candidate	NM_007275	1.11
hypothetical protein FLJ20694	NM_017928	1.11
eukaryotic translation initiation factor 3, subunit 10 (theta, 150/170kD)	AI123320	1.11
serine (or cysteine) proteinase inhibitor, clade B (ovalbumin), member 7	NM_003784	1.11
ELAV (embryonic lethal, abnormal vision, Drosophila)-like 1 (Hu antigen R)	BC003376	1.11
ABO blood group (transferase A, alpha 1-3-N-acetylgalactosaminyltransferase; transferase B, alpha 1-3-galactosyltransferase)	NM_020469	1.11
protein disulfide isomerase related protein (calcium-binding protein, intestinal-related)	BC006344	1.11
hypothetical protein FLJ22402	NM_024608	1.11
RGC32 protein	NM_014059	1.11
ubiquitination factor E4A (UFD2 homolog, yeast)	NM_004788	1.11
growth hormone inducible transmembrane protein	AL136713	1.11
lanosterol synthase (2,3-oxidosqualene-lanosterol cyclase)	D63807	1.11
protein with polyglutamine repeat; calcium (ca2+) homeostasis endoplasmic reticulum protein	NM_006387	1.11
dimethylarginine dimethylaminohydrolase 2	AI012008	1.11
F-box only protein 21	AB020682	1.11
KIAA0882 protein	BE646554	1.11

tyrosyl-DNA phodphodiesterase	NM_018319	1.11
actin related protein 2/3 complex, subunit 4 (20 kD)	BE891920	1.11
rab3 GTPase-activating protein, non-catalytic subunit (150kD)	NM_012414	1.11
group-specific component (vitamin D binding protein)	NM_000583	1.11
KIAA0494 gene product	AK001487	1.11
arginyl aminopeptidase (aminopeptidase B)	NM_020216	1.11
KIAA0692 protein	AW274877	1.11
fragile X mental retardation 2	AI001550	1.11
pre-mRNA splicing factor 17	BG528818	1.11
protein tyrosine phosphatase type IVA, member 1	AW165960	1.11
KIAA1638 protein	NM_025132	1.11
tubulin, beta, 2	AA515698	1.11
intercellular adhesion molecule 1 (CD54), human rhinovirus receptor	NM_000201	1.11
cAMP responsive element binding protein 1	AA161486	1.11
CGI-30 protein	NM_015958	1.11
diacylglycerol kinase, iota	NM_004717	1.11
Consensus includes gb:BG536224 /FEA=EST /DB_XREF=gi:13527769 /DB_XREF=est:602565445F1 /CLONE=IMAGE:4690258 /UG=Hs.123030 Human kappa-immunoglobulin germline pseu	BG536224	1.11
glutamate receptor, ionotropic, N-methyl D-aspartate 2A	NM_000833	1.11
glutamate decarboxylase 2 (pancreatic islets and brain, 65kD)	NM_000818	1.11
PBX/knotted 1 homeobox 1	NM_004571	1.11
dystonia 1, torsion (autosomal dominant; torsin A)	NM_000113	1.11
LIM domain only 6	U93305	1.11
cystatin F (leukocystatin)	AF031824	1.11
hypothetical protein FLJ22378	NM_025078	1.11
hypothetical protein F23149_1	NM_019088	1.11
inhibitor of growth family, member 3	NM_019071	1.11
zinc finger protein 278	AF254083	1.11
hypothetical protein FLJ20059	AW006750	1.11
muskelin 1, intracellular mediator containing kelch motifs	NM_013255	1.11
KIAA0301 protein	BE670928	1.11
cytochrome P450, subfamily IIA (phenobarbital-inducible), polypeptide 7, pseudogene 1	U22030	1.11
hypothetical protein FLJ20552	BG420893	1.11
hypothetical protein FLJ20506	NM_017848	1.11
Consensus includes gb:AA156865 /FEA=EST /DB_XREF=gi:1728480 /DB_XREF=est:zl20f08.s1 /CLONE=IMAGE:502503 /UG=Hs.22370 Homo sapiens mRNA; cDNA DKFZp56400122 (from	AA156865	1.11
Consensus includes gb:BE738425 /FEA=EST /DB_XREF=gi:10152417 /DB_XREF=est:601572441T1 /CLONE=IMAGE:3839147 /UG=Hs.241507 ribosomal protein S6	BE738425	1.11
nucleosome assembly protein 1-like 1	AI985751	1.11
KIAA0699 protein	AI934125	1.11
squamous cell carcinoma antigen recognised by T cells 3	D63879	1.11
hypothetical protein FLJ21432	NM_024551	1.11
AD024 protein	AF225416	1.11
hypothetical protein FLJ20984	NM_024630	1.10
vitamin D (1,25- dihydroxyvitamin D3) receptor	NM_000376	1.10
acylphosphatase 1, erythrocyte (common) type	NM_001107	1.10
ferritin, light polypeptide	BF312331	1.10
Consensus includes gb:AL109622 /DEF=Human DNA sequence from clone RP3-526F5 on chromosome Xq26.3-28. Contains the gene for a novel protein similar to UBE2N (ubiquitin-conjugat	AL109622	1.10
hypothetical protein DKFZp56481162	NM_031305	1.10
apoptosis inhibitor	NM_021631	1.10
nuclear transcription factor Y, beta	AI804118	1.10
cholinergic receptor, nicotinic, gamma polypeptide	NM_005199	1.10
Consensus includes gb:AV694312 /FEA=EST /DB_XREF=gi:10296175 /DB_XREF=est:AV694312 /CLONE=GKCELG01 /UG=Hs.250638 Homo sapiens mRNA full length insert cDNA clone EUR	AV694312	1.10
serine palmitoyltransferase, long chain base subunit 1	AL568804	1.10
karyopherin alpha 6 (importin alpha 7)	AK002111	1.10
neuronatin	NM_005386	1.10
WD40 repeat domain 11 protein	NM_018117	1.10
ATPase, Cu++ transporting, alpha polypeptide (Menkes syndrome)	BE567813	1.10
KIAA0395 protein	AW292783	1.10
hypothetical protein FLJ11132	BC006222	1.10
KIAA0552 gene product	NM_014731	1.10
hypothetical protein MGC10986	NM_030576	1.10
protein kinase C, epsilon	NM_005400	1.10
PRO1848 protein	AF118078	1.10
Consensus includes gb:BF967998 /FEA=EST /DB_XREF=gi:12335213 /DB_XREF=est:602269506F1 /CLONE=IMAGE:4357777 /UG=Hs.21851 Homo sapiens cDNA FLJ12900 fis, clone NT2R	BF967998	1.10
sec13-like protein	AV701173	1.10

PPAR binding protein	NM_004774	1.10
laminin, alpha 5	BC003355	1.10
hypothetical protein	NM_019057	1.10
SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily a, member 4	AI684141	1.10
ceroid-lipofuscinosis, neuronal 3, juvenile (Batten, Spielmeyer-Vogt disease)	AC002544	1.10
NCK protein	BC006356	1.10
Consensus includes gb:AW593996 /FEA=EST /DB_XREF=gi:7281254 /DB_XREF=est:hg41g06.x1 /CLONE=IMAGE:2948218 /UG=Hs.124969 Homo sapiens clone 24707 mRNA sequence	AW593996	1.10
Consensus includes gb:BG491393 /FEA=EST /DB_XREF=gi:13452905 /DB_XREF=est:602535682F1 /CLONE=IMAGE:4684748 /UG=Hs.183110 ESTs, Weakly similar to ALUC_HUMAN !!!! AL	BG491393	1.10
retinoid X receptor, beta	BF337038	1.10
lysosomal-associated membrane protein 2	NM_002294	1.10
albumin	AF116645	1.10
mitogen-activated protein kinase kinase kinase kinase 5	AW298170	1.10
maternally expressed 3	AF119863	1.10
3-oxoacid CoA transferase	NM_000436	1.10
ninjurin 2	NM_016533	1.10
similar to yeast BET3 (S. cerevisiae)	NM_014408	1.10
toll-like receptor 1	AL050262	1.10
retinoic acid repressible protein	AW188134	1.10
Cdc42 guanine exchange factor (GEF) 9	NM_015185	1.10
receptor-interacting serine-threonine kinase 2	AF064824	1.10
Meis1, myeloid ecotropic viral integration site 1 homolog 2 (mouse)	NM_020149	1.10
heme-regulated initiation factor 2-alpha kinase	AW007368	1.10
proline rich 1	NM_021225	1.10
Consensus includes gb:AL043266 /FEA=EST /DB_XREF=gi:5935844 /DB_XREF=est:DKFZp434L1423_s1 /CLONE=DKFZp434L1423 /UG=Hs.111334 ferritin, light polypeptide	AL043266	1.10
azotrophin	BC003404	1.10
cAMP responsive element modulator	U44836	1.10
chromogranin A (parathyroid secretory protein 1)	NM_001275	1.10
gamma-tubulin complex protein 2	AA016035	1.10
solute carrier family 25 (mitochondrial carrier; peroxisomal membrane protein, 34kD), member 17	BC005957	1.10
hypothetical protein FLJ11011	NM_018299	1.10
glycogenin	AF087942	1.10
heterogeneous nuclear ribonucleoprotein H2 (H')	NM_019597	1.10
hypothetical protein FLJ10420	NM_018090	1.10
hypothetical protein FLJ13646	NM_024584	1.10
Consensus includes gb:AF070587.1 /DEF=Homo sapiens clone 24741 mRNA sequence. /FEA=mRNA /DB_XREF=gi:3387960 /UG=Hs.25770 Homo sapiens clone 24741 mRNA sequence	AF070587	1.10
KIAA0923 protein	RS2678	1.10
Consensus includes gb:BF062139 /FEA=EST /DB_XREF=gi:10821049 /DB_XREF=est:7k72f10.x1 /CLONE=IMAGE:3481099 /UG=Hs.282387 polymerase (RNA) III (DNA directed) (32kD) /F	BF062139	1.10
gb:NM_018611.1 /DEF=Homo sapiens hypothetical protein PRO1966 (PRO1966), mRNA. /FEA=mRNA /GEN=PRO1966 /PROD=hypothetical protein PRO1966 /DB_XREF=gi:8924101 /UG=	NM_018611	1.10
peroxisomal biogenesis factor 6	NM_000287	1.10
chromosome 5 open reading frame 5	NM_016603	1.10
solute carrier family 19 (folate transporter), member 1	U15939	1.10
CGI-49 protein	NM_016002	1.10
acrosomal vesicle protein 1	NM_020109	1.10
ral guanine nucleotide dissociation stimulator	AF295773	1.10
Consensus includes gb:AA676803 /FEA=EST /DB_XREF=gi:2657325 /DB_XREF=est:zj65b04.s1 /CLONE=IMAGE:455119 /UG=Hs.13996 Homo sapiens cDNA: FLJ23260 fis, clone COL05804	AA676803	1.10
chromosome 21 open reading frame 66	NM_013329	1.10
arylalkylamine N-acetyltransferase	NM_001088	1.10
MINA (never in mitosis gene a)-related kinase 1	AI936517	1.10
kinesin-like 7	NM_020242	1.10
supertiller viralicidic activity 2-like (S. cerevisiae)	NM_006929	1.10
CLIP-associating protein 2	AL137636	1.10
heterogeneous nuclear ribonucleoprotein D (AU-rich element RNA binding protein 1, 37kD)	D55672	1.10
Consensus includes gb:AW194543 /FEA=EST /DB_XREF=gi:6473381 /DB_XREF=est:xb27c02.x1 /CLONE=IMAGE:2577506 /UG=Hs.234573 Homo sapiens mRNA for TL132	AW194543	1.10
nuclear receptor coactivator 3	AI761748	1.10
prostate epithelium-specific Ets transcription factor	AI435670	1.10
DKFZP586G011 protein	AK023204	1.10
cylindromatosis (turban tumor syndrome)	AK024212	1.10
peroxisomal biogenesis factor 16	AA523441	1.10
Consensus includes gb:H53689 /FEA=EST /DB_XREF=gi:993836 /DB_XREF=est:yu38g07.r1 /CLONE=IMAGE:236124 /UG=Hs.306239 Homo sapiens clone ASPBLL54 immunoglobulin lambd	H53689	1.10
golgi phosphoprotein 3 (coat-protein)	NM_022130	1.10
nucleolar protein (KKE/D repeat)	NM_006392	1.10
baculoviral IAP repeat-containing 1	AI817801	1.10
myotubularin related protein 6	U47635	1.10

MRE11 meiotic recombination 11 homolog A (S. cerevisiae)	NM_005590	1.10
ankyrin 1, erythrocytic	NM_020479	1.10
sorting nexin 11	NM_013323	1.10
mucolipin 1	NM_020533	1.10
nuclear LIM interactor-interacting factor	NM_021198	1.10
meningioma expressed antigen 6 (coiled-coil proline-rich)	NM_005930	1.10
hypothetical protein FLJ20080	NM_017657	1.10
cytochrome P450, subfamily XXVIIB (25-hydroxyvitamin D-1-alpha-hydroxylase), polypeptide 1	NM_000785	1.10
hypothetical protein FLJ10415	NM_018089	1.10
dendritic cell protein	NM_006360	1.10
hypothetical protein IMAGE:4215339	AW182303	1.10
hypothetical protein FLJ14213	NM_024841	1.10
RAB27A, member RAS oncogene family	AF125393	1.10
1-acylglycerol-3-phosphate O-acyltransferase 2 (lysophosphatidic acid acyltransferase, beta)	U56418	1.10
hypothetical protein FLJ22418	NM_024626	1.10
chromosome 21 open reading frame 2	U84569	1.10
cytochrome P450, 51 (lanosterol 14-alpha-demethylase)	NM_000786	1.10
hypothetical protein MGC3265	NM_024028	1.10
Consensus includes gb:U80770.1 /DEF=Human EST clone 251800 mariner transposon Hsmar1 sequence. /FEA=mRNA /DB_XREF=gi:2921608 /UG=Hs.42192 Human EST clone 251800 ma	U80770	1.10
serine/threonine kinase 4	NM_006282	1.10
CamKI-like protein kinase	NM_020397	1.10
small inducible cytokine A1, I-309	NM_002981	1.10
transcriptional adaptor 2 (ADA2 homolog, yeast)-like	AF064094	1.10
sema domain, immunoglobulin domain (Ig), short basic domain, secreted, (semaphorin) 3F	U38276	1.10
COP9 subunit 6 (MOV34 homolog, 34 kD)	W63732	1.10
endothelin 2	NM_001956	1.10
Rho guanine exchange factor (GEF) 11	AI090007	1.10
Consensus includes gb:AI378788 /FEA=EST /DB_XREF=gi:4188641 /DB_XREF=est:tc67b04.x1 /CLONE=IMAGE:2069647 /UG=Hs.153445 Human mRNA for unknown product, partial cds	AI378788	1.10
kinesin-like 1	NM_004523	1.10
dolichyl-phosphate mannosyltransferase polypeptide 2, regulatory subunit	AF061729	1.10
Consensus includes gb:BF982927 /FEA=EST /DB_XREF=gi:12385739 /DB_XREF=est:602306318F1 /CLONE=IMAGE:4397525 /UG=Hs.324787 solute carrier family 5 (inositol transporters)	BF982927	1.10
serine/threonine kinase 10	NM_005990	1.10
testis expressed sequence 13A	NM_031274	1.10
nuclear receptor coactivator 3	NM_006534	1.10
chromodomain helicase DNA binding protein 4	AI761771	1.10
SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily a, member 2	AW131754	1.10
RNA binding motif protein 12	AL514547	1.10
histone H2A F/Z variant	AA534860	1.10
rTS beta protein	NM_017512	1.10
sin3-associated polypeptide, 18kD	AW274856	1.10
aldehyde dehydrogenase 3 family, member B2	AA071510	1.10
ubiquitin specific protease 20	NM_006676	1.10
hypothetical protein MGC14797	AI816243	1.10
MHC class I polypeptide-related sequence A	NM_000247	1.10
U6 snRNA-associated Sm-like protein	AU153477	1.10
KIAA1668 protein	W46406	1.09
gb:L14454.1 /DEF=Human Ig rearranged mu-chain gene V-N-D-N-J-region, complete cds. /FEA=mRNA /DB_XREF=gi:398495 /FL=gb:L14454.1	L14454	1.09
SAM domain and HD domain, 1	NM_015474	1.09
solute carrier family 4, sodium bicarbonate cotransporter, member 7	AF053755	1.09
heat shock 70kD protein 9B (mortalin-2)	AA927701	1.09
Xq28, 2000bp sequence contg. ORF	AF319573	1.09
microtubule-associated protein, RP/EB family, member 2	NM_014268	1.09
KIAA0244 protein	NM_015153	1.09
dual specificity phosphatase 2	NM_004418	1.09
hypothetical protein FLJ10856	NM_018247	1.09
zinc finger protein 211	NM_006385	1.09
sialoadhesin	N53555	1.09
methylcrotonoyl-Coenzyme A carboxylase 2 (beta)	AW439494	1.09
ubiquitin-like 4	BE543027	1.09
DKFZP566D213 protein	NM_015513	1.09
neuroblastoma, suppression of tumorigenicity 1	NM_005380	1.09
c6.1A	NM_024332	1.09
hypothetical protein FLJ23529	AK027182	1.09

prosaposin (variant Gaucher disease and variant metachromatic leukodystrophy)	M32221	1.09
BCL2/adenovirus E1B 19kD interacting protein 3-like	AF060922	1.09
tripartite motif-containing 17	NM_016102	1.09
retinoic acid induced 15	U50383	1.09
hypothetical protein FLJ20627	NM_017909	1.09
6 antigen, family B, 1 (prostate associated)	NM_003785	1.09
vesicle-associated membrane protein 1 (synaptobrevin 1)	NM_016830	1.09
gb:NM_001222.1 /DEF=Homo sapiens calciumcalmodulin-dependent protein kinase (CaM kinase) II gamma (CAMK2G), mRNA. /FEA=mRNA /GEN=CAMK2G /PROD=calciumcalmodulin-depen	NM_001222	1.09
hypothetical protein	BF572868	1.09
leucine carboxyl methyltransferase	BC001214	1.09
SEC63 protein	NM_007214	1.09
regulator of G-protein signalling 20	AF074979	1.09
z 003 protein	NM_020232	1.09
SRY (sex determining region Y)-box 22	NM_006943	1.09
Wiskott-Aldrich syndrome protein interacting protein	NM_003387	1.09
putatative 28 kDa protein	NM_020143	1.09
hypothetical protein FLJ21562	NM_025113	1.09
hypothetical protein FLJ23322	NM_024955	1.09
density-regulated protein	AB014731	1.09
chondroitin sulfate proteoglycan 6 (bamacan)	AJ373676	1.09
hypothetical protein MGC11349	NM_025112	1.09
poly(rC) binding protein 2	AW103422	1.09
hypothetical protein MGC13523	AL049699	1.09
zinc finger protein 277	NM_021994	1.09
matrix metalloproteinase 24 (membrane-inserted)	W45551	1.09
suppressor of var1, 3-like 1 (S. cerevisiae)	NM_003171	1.09
KIAA0409 protein	AB007869	1.09
small inducible cytokine subfamily E, member 1 (endothelial monocyte-activating)	BF589679	1.09
amyotrophic lateral sclerosis 2 (juvenile) chromosome region, candidate 3	NM_015049	1.09
CGI-02 protein	AI249752	1.09
joined to JAZF1	BF382924	1.09
BH-protocadherin (brain-heart)	NM_002589	1.09
DiGeorge syndrome critical region gene 6 like	AA631156	1.09
hypothetical protein FLJ10057	NM_017984	1.09
KIAA1111 protein	AW249934	1.09
NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 4 (15kD, B15)	NM_004547	1.09
active BCR-related gene	L19704	1.09
RNA polymerase I 16 kDa subunit	NM_015972	1.09
galactose-4-epimerase, UDP-	AL031295	1.09
hypothetical protein FLJ23441	NM_024678	1.09
Sp2 transcription factor	D28588	1.09
heterogeneous nuclear ribonucleoprotein H3 (2H9)	NM_021644	1.09
CD1D antigen, d polypeptide	NM_001766	1.09
Consensus includes gb:AL023584 /DEF=Human DNA sequence from clone 67K17 on chromosome 6q24.1-24.3. Contains the HIVEP2 (Schnurri-2) gene for HIV type 1 Enhancer-binding Prot	AL023584	1.09
KIAA0758 protein	N95226	1.09
glucagon receptor	U03469	1.09
19A24 protein	NM_021181	1.09
optic atrophy 3 (autosomal recessive, with chorea and spastic paraplegia)	NM_025136	1.09
RAB3 GTPase-ACTIVATING PROTEIN	AK022494	1.09
hypothetical protein FLJ11585	NM_023075	1.09
F-box only protein 2	NM_012168	1.09
hypothetical protein MGC5576	NM_024056	1.09
POU domain, class 5, transcription factor 1	AF268615	1.09
high-mobility group 20B	N25429	1.09
Consensus includes gb:AI017564 /FEA=EST /DB_XREF=gi:3231900 /DB_XREF=est:ou23e03.x1 /CLONE=IMAGE:1627132 /UG=Hs.95631 Human normal keratinocyte mRNA	AI017564	1.09
catechol-O-methyltransferase	BC000419	1.09
DKFZP564G0222 protein	NM_014044	1.09
hypothetical protein FLJ20260	AF323729	1.09
engulfment and cell motility 2 (ced-12 homolog, C. elegans)	NM_022086	1.09
H1 histone family, member 2	BC002649	1.09
ets variant gene 5 (ets-related molecule)	X76184	1.09
G protein-coupled receptor 63	NM_030784	1.09
hypothetical gene MGC19595	BC003700	1.09

NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 1 (7kD, MNLL)	NM_004545	1.09
DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 34	AI924903	1.09
mitogen-activated protein kinase 7	U29725	1.09
ubiquitin specific protease 9, X chromosome (Drosophila fat facets related)	AA824386	1.09
trophinin	AF349720	1.09
retinoblastoma binding protein 7	NM_002893	1.09
thrombopoietin (myeloproliferative leukemia virus oncogene ligand, megakaryocyte growth and development factor)	D32047	1.09
EphA1	Z27409	1.09
hypothetical protein FLJ20345	NM_017777	1.09
signal peptidase complex (18kD)	NM_014300	1.09
hypothetical protein FLJ21820	NM_021925	1.09
hypothetical protein FLJ23514	NM_021827	1.09
hypothetical protein MGC5149	U79260	1.09
DKFZP564I052 protein	AL080063	1.09
similar to yeast Upf3, variant A	NM_023011	1.09
diptheria toxin resistance protein required for diphthamide biosynthesis (Saccharomyces)-like 1	NM_001383	1.09
interleukin 5 receptor, alpha	NM_000564	1.09
Consensus includes gb:AK022632.1 /DEF=Homo sapiens cDNA FLJ12570 fis, clone NT2RM4000895. /FEA=mRNA /DB_XREF=gi:10434135 /UG=Hs.142076 Homo sapiens cDNA FLJ12570 f	AK022632	1.09
KIAA0648 protein	AK021757	1.09
HSPC052 protein	NM_014150	1.09
zinc finger protein 36, C3H type-like 2	NM_006887	1.09
pannexin 1	NM_015368	1.09
Rho GTPase activating protein 4	NM_001666	1.09
coagulation factor XI (plasma thromboplastin antecedent)	NM_000128	1.09
beaded filament structural protein 2, phakinin	NM_003571	1.09
acetyl-Coenzyme A acyltransferase 2 (mitochondrial 3-oxoacyl-Coenzyme A thiolase)	AW072302	1.09
Cluster Incl. AW014299:UI-H-BI0p-aas-a-07-0-UI.s1 Homo sapiens cDNA, 3' end /clone=IMAGE-2710573 /clone_end=3' /gb=AW014299 /gi=5863056 /ug=Hs.238968 /len=391	AW014299	1.09
ubiquitin-conjugating enzyme E2D 2 (UBC4/5 homolog, yeast)	NM_003339	1.09
CD28 antigen (Tp44)	NM_006139	1.09
ret finger protein 2	NM_005798	1.09
regulatory factor X, 1 (influences HLA class II expression)	NM_002918	1.09
interferon-related developmental regulator 1	AA747426	1.09
BCL2-like 2	D87461	1.09
gb:AF277175.1 /DEF=Homo sapiens PNAS-138 mRNA, complete cds. /FEA=mRNA /PROD=PNAS-138 /DB_XREF=gi:12751080 /UG=Hs.326790 Homo sapiens PNAS-138 mRNA, complete cd	AF277175	1.09
BRCA1 associated protein	AF035620	1.09
Rho GTPase activating protein 8	NM_015366	1.09
bone morphogenetic protein 15	NM_005448	1.09
Lsm3 protein	NM_014463	1.09
cell division cycle 25B	AL109804	1.09
adenylate kinase 1	NM_000476	1.09
palmitoyl-protein thioesterase 2	AF020544	1.09
bone morphogenetic protein 3 (osteogenic)	NM_001201	1.09
ribonuclease 6 precursor	NM_003730	1.09
KIAA0411 gene product	AB007871	1.09
dual-specificity tyrosine-(Y)-phosphorylation regulated kinase 2	Y09216	1.09
matrix metalloproteinase 14 (membrane-inserted)	X83535	1.09
hypothetical protein MGC2574	NM_024098	1.09
opioid receptor, kappa 1	NM_000912	1.09
chorea acanthocytosis	AB023203	1.09
5-methyltetrahydrofolate-homocysteine methyltransferase reductase	N29717	1.09
DKFZP586D2223 protein	AK023092	1.09
H2A histone family, member P	NM_021064	1.09
KIAA1139 protein	AK025262	1.09
centaurin, beta 2	D26069	1.09
KIAA1354 protein	AW138594	1.09
hypothetical protein FLJ22761	NM_025130	1.09
hypothetical protein FLJ20258	BC004907	1.09
ring finger protein 13	NM_007282	1.09
nuclear receptor subfamily 2, group F, member 6	AI935343	1.09
fibroblast growth factor receptor 4	NM_002011	1.09
protein phosphatase 2 (formerly 2A), regulatory subunit B" (PR 72), alpha isoform and (PR 130), beta isoform	AL389975	1.09
dishevelled, dsh homolog 3 (Drosophila)	U49262	1.09
PAI-1 mRNA-binding protein	BC002488	1.09

hypothetical protein MGC12335	AL022724	1.09
homeo box A1	S79910	1.09
ER to nucleus signalling 1	NM_001433	1.09
KIAA1698 protein	BF446693	1.09
NAD kinase	BC001709	1.08
KIAA0783 gene product	NM_014660	1.08
core-binding factor, runt domain, alpha subunit 2; translocated to, 1; cyclin D-related	X79990	1.08
Consensus includes gb:AK000864.1 /DEF=Homo sapiens cDNA FLJ10002 fis, clone HEMBA1000046. /FEA=mRNA /DB_XREF=gi:7021188 /UG=Hs.296522 Homo sapiens cDNA FLJ10002 fis	AK000864	1.08
Consensus includes gb:R61519 /FEA=EST /DB_XREF=gi:832214 /DB_XREF=est:yh16c04.s1 /CLONE=IMAGE:37896 /UG=Hs.185807 Homo sapiens clone 24758 mRNA sequence	R61519	1.08
DKFZP586C1620 protein	AV694093	1.08
small nuclear RNA activating complex, polypeptide 4, 190kD	NM_003086	1.08
v-maf musculoaponeurotic fibrosarcoma oncogene homolog K (avian)	NM_002360	1.08
SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily a, member 5	NM_003601	1.08
FGF receptor activating protein 1	AL049261	1.08
HERV-H LTR-associating 2	NM_007072	1.08
ubiquinol-cytochrome c reductase binding protein	BC005230	1.08
Consensus includes gb:AC005756 /DEF=Homo sapiens chromosome 19, fosmid 39347 /FEA=CDS /DB_XREF=gi:3688091 /UG=Hs.247754 Homo sapiens chromosome 19, fosmid 39347	AC005756	1.08
wingless-type MMTV integration site family, member 16	NM_016087	1.08
HSPC041 protein	NM_016099	1.08
expressed in activated T/LAK lymphocytes	NM_007267	1.08
RNA binding motif protein 8A	AF182415	1.08
Consensus includes gb:AI267546 /FEA=EST /DB_XREF=gi:3886713 /DB_XREF=est:aq90f05.x1 /CLONE=IMAGE:2038209 /UG=Hs.90798 Human clone 23695 mRNA sequence	AI267546	1.08
capicua homolog (Drosophila)	AB002304	1.08
gb:Z25432.1 /DEF=H.sapiens protein-serinethreonine kinase gene, complete CDS. /FEA=mRNA /PROD=protein-serinethreonine kinase /DB_XREF=gi:405742 /FL=gb:Z25432.1	Z25432	1.08
calcium/calmodulin-dependent serine protein kinase (MAGUK family)	NM_003688	1.08
hypothetical protein FLJ00002	AK024446	1.08
KIAA1139 protein	AI970096	1.08
gb:M87268.1 /DEF=Human IgM VDJ-region mRNA, complete cds. /FEA=CDS /GEN=IGHM /PROD=IgM /DB_XREF=gi:186197 /FL=gb:M87268.1	M87268	1.08
pro-melanin-concentrating hormone	NM_002674	1.08
RAS-related on chromosome 22	NM_006477	1.08
ubiquilin 2	AK001029	1.08
ubiquitin-conjugating enzyme E2D 2 (UBC4/5 homolog, yeast)	BE621259	1.08
KIAA0431 protein	AI744148	1.08
KIAA0438 gene product	AA142966	1.08
hypothetical protein MGC4276 similar to CG8198	BG387555	1.08
DKFZP434C212 protein	AI797397	1.08
Consensus includes gb:AF308291.1 /DEF=Homo sapiens serologically defined breast cancer antigen NY-BR-46 mRNA, partial cds. /FEA=mRNA /PROD=serologically defined breast cancer a	AF308291	1.08
KIAA0904 protein	AW305119	1.08
mitogen-activated protein kinase kinase 5	U71087	1.08
Consensus includes gb:AK026155.1 /DEF=Homo sapiens cDNA: FLJ22502 fis, clone HRC11383. /FEA=mRNA /DB_XREF=gi:10438915 /UG=Hs.298140 Homo sapiens cDNA: FLJ22502 fis, cl	AK026155	1.08
neurofibromin 2 (bilateral acoustic neuroma)	AF122827	1.08
homeo box B7	AW102783	1.08
Cas-Br-M (murine) ectropic retroviral transforming sequence c	NM_012116	1.08
hypothetical protein FLJ22055	NM_024779	1.08
thioredoxin reductase beta	AF201385	1.08
Consensus includes gb:AL117381 /DEF=Human DNA sequence from clone RP5-857M17 on chromosome 20 Contains ESTs, STSs, GSSs and CpG islands. Contains the 3 part of the ID1 gene f	AL117381	1.08
E2F transcription factor 2	NM_004091	1.08
potassium large conductance calcium-activated channel, subfamily M beta member 3	NM_014407	1.08
calmodulin 1 (phosphorylase kinase, delta)	N25325	1.08
mitogen-activated protein kinase kinase 4	NM_003010	1.08
cytochrome P450, subfamily XXIV (vitamin D 24-hydroxylase)	NM_000782	1.08
engulfment and cell motility 2 (ced-12 homolog, C. elegans)	W22924	1.08
SRY (sex determining region Y)-box 2	L07335	1.08
S100 calcium binding protein A13	NM_005979	1.08
hexose-6-phosphate dehydrogenase (glucose 1-dehydrogenase)	NM_004285	1.08
KIAA0495	AB007964	1.08
branched chain keto acid dehydrogenase E1, alpha polypeptide (maple syrup urine disease)	NM_000709	1.08
gene from NF2/meningioma region of 22q12	AB023200	1.08
dermatopontin	NM_001937	1.08
hypothetical protein FLJ20241	NM_017721	1.08
hypothetical protein FLJ11730	NM_022756	1.08
pro-platelet basic protein (includes platelet basic protein, beta-thromboglobulin, connective tissue-activating peptide III, neutrophil-activating peptide-2)	R64130	1.08
lifeguard	NM_012306	1.08

inner centromere protein antigens (135kD, 155kD)	NM_020238	1.08
histamine receptor H3	AB045369	1.08
KIAA0082 protein	O43949	1.08
hypothetical protein FLJ10206	NM_018025	1.08
transcription factor 12 (HTF4, helix-loop-helix transcription factors 4)	AL559478	1.08
bile acid Coenzyme A: amino acid N-acyltransferase (glycine N-choloyltransferase)	NM_001701	1.08
BCL2/adenovirus E1B 19kD interacting protein 2	BC002461	1.08
hypothetical protein FLJ12567	NM_024714	1.08
immunoglobulin kappa variable 1-13	AW408194	1.08
ubiquitin specific protease 4 (proto-oncogene)	AI346043	1.08
mitogen-activated protein kinase kinase 3	AA780381	1.08
zinc finger protein 91 (HPF7, HTF10)	NM_003430	1.08
integrin, alpha 7	AK022548	1.08
glyceronephosphate O-acyltransferase	NM_014236	1.08
tumor protein, translationally-controlled 1	AL565449	1.08
serine racemase	NM_021947	1.08
sorbitol dehydrogenase	NM_003104	1.08
fatty-acid-Coenzyme A ligase, long-chain 3	D89053	1.08
aquaporin 3	N74607	1.08
SFRS protein kinase 2	AC005070	1.08
hypothetical protein FLJ13102	NM_024887	1.08
KIAA0700 protein	AB014600	1.08
growth arrest and DNA-damage-inducible, gamma	NM_006705	1.08
UDP-GlcNAc:betaGal beta-1,3-N-acetylglucosaminyltransferase 6	NM_006876	1.08
cadherin, EGF LAG seven-pass G-type receptor 1 (flamingo homolog, Drosophila)	AL031588	1.08
discoidin domain receptor family, member 1	U48705	1.08
hydroxysteroid (17-beta) dehydrogenase 4	NM_000414	1.08
hypothetical protein FLJ20030	BG498776	1.08
CGI-130 protein	BC001671	1.08
synovial sarcoma translocation gene on chromosome 18-like 2	NM_016305	1.08
Consensus includes gb:AF217536.1 /DEF=Homo sapiens truncated mevalonate kinase mRNA, partial cds, alternatively spliced. /FEA=mRNA /PROD=truncated mevalonate kinase /DB_XREF=	AF217536	1.08
Tax1 (human T-cell leukemia virus type I) binding protein 1	NM_006024	1.08
hypothetical protein FLJ22609	NM_022072	1.08
myosin, light polypeptide 4, alkali; atrial, embryonic	X52005	1.08
catenin (cadherin-associated protein), alpha-like 1	NM_003798	1.08
protein kinase D2	AF309082	1.08
artemin	AF120274	1.08
hypothetical protein FLJ22643	NM_024635	1.08
polymerase (DNA directed), gamma 2, accessory subunit	NM_007215	1.08
immunoglobulin superfamily, member 4	NM_014333	1.08
hypothetical protein MGC2550	NM_024071	1.08
KIAA0650 protein	AV699744	1.08
Consensus includes gb:AU147830 /FEA=EST /DB_XREF=gi:11009351 /DB_XREF=est:AU147830 /CLONE=MAMMA1001818 /UG=Hs.296713 Homo sapiens cDNA FLJ12295 fis, clone MAMM	AU147830	1.08
solute carrier family 21 (organic anion transporter), member 3	AF085224	1.08
collagen, type XIX, alpha 1	D38163	1.08
hypothetical protein DKFZp547M136 similar to widely-interspaced zinc finger motifs	AA422049	1.08
serine/arginine repetitive matrix 1	AU147713	1.08
neurotensin receptor 2	NM_012344	1.08
carnitine acetyltransferase	BC000723	1.08
Consensus includes gb:AI925734 /FEA=EST /DB_XREF=gi:5661698 /DB_XREF=est:wo34g08.x1 /CLONE=IMAGE:2457278 /UG=Hs.110613 KIAA0220 protein	AI925734	1.08
putative L-type neutral amino acid transporter	AU154782	1.08
protein phosphatase 1, regulatory (inhibitor) subunit 11	NM_021959	1.08
SP110 nuclear body protein	AF280094	1.08
proteasome (prosome, macropain) 26S subunit, non-ATPase, 10	NM_002814	1.08
HIV-1 rev binding protein 2	NM_007043	1.08
neural cell adhesion molecule 1	U63041	1.08
endoplasmic reticulum oxidoreductin 1-Lbeta	NM_019891	1.08
G protein-coupled receptor 44	AF118265	1.08
retinoic acid receptor, alpha	T91506	1.08
heparan sulfate (glucosamine) 3-O-sulfotransferase 3B1	NM_006041	1.08
rasA binding protein 1	NM_006788	1.08
interferon regulatory factor 5	NM_002200	1.08
colony stimulating factor 1 (macrophage)	M37435	1.08

interleukin 9 receptor	NM_002186	1.08
butyrophilin, subfamily 2, member A2	NM_006995	1.08
type I transmembrane receptor (seizure-related protein)	NM_012410	1.08
hypothetical protein DKFZp547M136 similar to widely-interspaced zinc finger motifs	AI828531	1.08
Consensus includes gb:AU134977 /FEA=EST /DB_XREF=gi:10995516 /DB_XREF=est:AU134977 /CLONE=PLACE1000926 /UG=Hs.322149 Human clone 137308 mRNA, partial cds	AU134977	1.08
KIAA1053 protein	AL117523	1.08
tripartite motif-containing 2	AW071795	1.08
transducer of ERBB2, 1	AA675892	1.08
COX17 homolog, cytochrome c oxidase assembly protein (yeast)	NM_005694	1.08
hypothetical protein FLJ22167	AA401703	1.08
hypothetical protein FLJ12707	NM_022067	1.08
solute carrier family 7, (cationic amino acid transporter, y+ system) member 11	AI040875	1.08
hypothetical protein PRO0899	NM_018565	1.08
D site of albumin promoter (albumin D-box) binding protein	U79283	1.07
hypothetical protein	NM_023070	1.07
hypothetical protein FLJ20030	NM_017627	1.07
calmegin	NM_004362	1.07
MAX protein	BC003525	1.07
KIAA0368 protein	AI002366	1.07
dihydrolipoamide branched chain transacylase (E2 component of branched chain keto acid dehydrogenase complex; maple syrup urine disease)	M27093	1.07
activin A receptor, type IB	NM_020328	1.07
poly(A)-specific ribonuclease (deadenylation nuclease)	NM_002582	1.07
Consensus includes gb:AA205593 /FEA=EST /DB_XREF=gi:1803601 /DB_XREF=est:zq68c06.s1 /CLONE=IMAGE:646762 /UG=Hs.26837 Homo sapiens mRNA; cDNA DKFZp586K1123 (from	AA205593	1.07
hypothetical protein DKFZp564D0478	AI828080	1.07
ubiquitin specific protease 5 (isopeptidase T)	NM_003481	1.07
deoxyhypusine synthase	NM_013407	1.07
partial transcript encompassing THC211630 gene	AI010070	1.07
6 protein-coupled receptor kinase-interactor 1	NM_014030	1.07
Hr44 antigen	X91103	1.07
apoptosis inhibitor 5	AF229253	1.07
hypothetical protein MGC3262	Z78349	1.07
retinoic acid repressible protein	NM_016167	1.07
Cluster Incl. N92708:zb44c04.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-306438 /clone_end=3 /gb=N92708 /gi=1265017 /ug=Hs.32374 /len=668	N92708	1.07
transforming growth factor beta-activated kinase-binding protein 1	NM_006116	1.07
major histocompatibility complex, class II, DO beta	NM_002120	1.07
hypothetical protein FLJ14639	AI884627	1.07
Consensus includes gb:W74494 /FEA=EST /DB_XREF=gi:1384781 /DB_XREF=est:zd75l10.s1 /CLONE=IMAGE:346507 /UG=Hs.284275 Homo sapiens PAK2 mRNA, complete cds /FL=gb:AF	W74494	1.07
Consensus includes gb:AW979196 /FEA=EST /DB_XREF=gi:8170484 /DB_XREF=est:EST391306 /UG=Hs.292713 ESTs, Moderately similar to ALU1_HUMAN ALU SUBFAMILY J SEQUENCE CO	AW979196	1.07
hypothetical protein FLJ23393	NM_024639	1.07
trinucleotide repeat containing 11 (THR-associated protein, 230 kDa subunit)	AF132033	1.07
nuclear protein, ataxia-telangiectasia locus	D83243	1.07
v-ets erythroblastosis virus E26 oncogene homolog 1 (avian)	NM_005238	1.07
potassium channel, subfamily K, member 10 (TREK-2)	NM_021161	1.07
immunoglobulin kappa constant	AF103574	1.07
HDCMA18P protein	AA160181	1.07
hypothetical protein FLJ11467	NM_024963	1.07
hypothetical protein FLJ20123	NM_017674	1.07
Consensus includes gb:AL109705.1 /DEF=Homo sapiens mRNA full length insert cDNA clone EUROIMAGE 73337. /FEA=mRNA /DB_XREF=gi:5689834 /UG=Hs.9997 Homo sapiens mRNA ful	AL109705	1.07
chondroitin sulfate proteoglycan 6 (bamacan)	AF020043	1.07
hypothetical protein PRO2325	AW170549	1.07
hypothetical protein FLJ20596	AL512707	1.07
hypoxia-inducible protein 2	NM_013332	1.07
heparan sulfate 2-O-sulfotransferase	NM_012262	1.07
HIV-1 Rev binding protein	AI742626	1.07
cannabinoid receptor 1 (brain)	NM_016083	1.07
ribosomal protein L14	BC000606	1.07
Consensus includes gb:AA071454 /FEA=EST /DB_XREF=gi:1578817 /DB_XREF=est:zm73c03.s1 /CLONE=IMAGE:531268 /UG=Hs.258725 ESTs, Moderately similar to ATPO_HUMAN ATP SY	AA071454	1.07
KIAA0237 gene product	BC003103	1.07
protein phosphatase 2, regulatory subunit B (B56), delta isoform	NM_006245	1.07
ATP/GTP-binding protein	NM_006831	1.07
polymerase (RNA) II (DNA directed) polypeptide B (140kD)	NM_000938	1.07
KIAA1067 protein	AB028990	1.07
KIAA0276 protein	D87466	1.07

Consensus includes gb:AI867198 /FEA=EST /DB_XREF=gi:5540214 /DB_XREF=est:wa01c11.x1 /CLONE=IMAGE:2296820 /UG=Hs.324787 solute carrier family 5 (inositol transporters), me	AI867198	1.07
interleukin 1 receptor, type II	U64094	1.07
Consensus includes gb:AK025343.1 /DEF=Homo sapiens cDNA: FLJ21690 fis, clone COL09538. /FEA=mRNA /DB_XREF=gi:10437841 /UG=Hs.306802 Homo sapiens cDNA: FLJ21690 fis, cl	AK025343	1.07
DAZ associated protein 2	NM_014764	1.07
hypothetical protein FLJ13798	NM_024773	1.07
transketolase-like 1	Z49258	1.07
holocarboxylase synthetase (biotin-[propionyl-Coenzyme A-carboxylase (ATP-hydrolysing)] ligase)	NM_000411	1.07
tumor necrosis factor receptor superfamily, member 10b	AF016266	1.07
vesicle-associated membrane protein 3 (cellubrevin)	BC003570	1.07
hypothetical protein FLJ23209	NM_024895	1.07
endothelin receptor type A	S81545	1.07
KIAA0471 gene product	BG107203	1.07
2',5'-oligoadenylate synthetase 1 (40-46 kD)	NM_016816	1.07
small nuclear RNA activating complex, polypeptide 3, 50kD	NM_003084	1.07
tripartite motif-containing 15	AF220133	1.07
40S ribosomal protein S27 isoform	NM_015920	1.07
Consensus includes gb:AL137784 /DEF=Human DNA sequence from clone RP1-199I3 on chromosome 6 Contains ESTs, STSs, GSSs and CpG islands. Contains part of the gene for a novel pro	AL137784	1.07
serine/threonine kinase 17a (apoptosis-inducing)	AW194730	1.07
mitogen-activated protein kinase 11	U53442	1.07
hypothetical protein MGC10765	NM_024345	1.07
adenomatous polyposis coli like	AL078616	1.07
mature T-cell proliferation 1	Z24459	1.07
adducin 3 (gamma)	AI763123	1.07
PR domain containing 2, with ZNF domain	AI681013	1.07
dystrophin (muscular dystrophy, Duchenne and Becker types), includes DXS142, DXS164, DXS206, DXS230, DXS239, DXS268, DXS269, DXS270, DXS272	NM_004010	1.07
phosphatidic acid phosphatase type 2C	AF047760	1.07
sarcosine dehydrogenase	AF047004	1.07
uncoupling protein 2 (mitochondrial, proton carrier)	U94592	1.07
ubiquitin-activating enzyme E1-like	L13852	1.07
KIAA0239 protein	AI735639	1.07
CGI-04 protein	NM_015936	1.07
Hsp90-associating relative of Cdc37	NM_017913	1.07
RAB14, member RAS oncogene family	AF112206	1.07
stanniocalcin 2	AI435828	1.07
A kinase (PRKA) anchor protein 1	U34074	1.07
hepatocyte growth factor (hepapoietin A; scatter factor)	M60718	1.07
inorganic pyrophosphatase	NM_006903	1.07
glutamate receptor, ionotrophic, AMPA 3	NM_007325	1.07
protein kinase C binding protein 1	AB032951	1.07
ribonuclease P (30kD)	NM_006413	1.07
H2A histone family, member G	NM_021065	1.07
gb:NM_015961.1 /DEF=Homo sapiens CGI-34 protein (LOC51612), mRNA. /FEA=mRNA /GEN=LOC51612 /PROD=CGI-34 protein /DB_XREF=gi:7706273 /UG=Hs.36237 CGI-34 protein /FL=	NM_015961	1.07
tankyrase, TRF1-interacting ankyrin-related ADP-ribose polymerase	AF070613	1.07
nuclear receptor subfamily 3, group C, member 1	X03348	1.07
monocyte to macrophage differentiation-associated	NM_012329	1.07
KIAA1069 protein	AK022610	1.07
zinc finger protein 117 (HPF9)	NM_024498	1.07
hypothetical protein FLJ10842	AI278150	1.07
gb:M17565.1 /DEF=Human MHC class II DQ-beta associated with DRw6, DQw1 protein, complete cds. /FEA=mRNA /GEN=HLA-DQB1 /DB_XREF=gi:188188 /FL=gb:M17565.1	M17565	1.07
inositol 1,4,5-triphosphate receptor, type 2	AI963873	1.07
chloride intracellular channel 1	U85978	1.07
ABO blood group (transferase A, alpha 1-3-N-acetylgalactosaminyltransferase; transferase B, alpha 1-3-galactosyltransferase)	U15197	1.07
protein kinase C, theta	L01087	1.07
peptidylglycine alpha-amidating monooxygenase	AI022882	1.07
zinc finger protein 197	NM_006991	1.07
Ras association (RalGDS/AF-6) domain family 2	NM_014737	1.07
ubiquitin specific protease 9, X chromosome (Drosophila fat facets related)	NM_004652	1.07
hypothetical protein MGC4707	NM_024113	1.07
mesoderm specific transcript homolog (mouse)	NM_002402	1.07
gb:AF130050.1 /DEF=Homo sapiens clone FLB3442 PRO0872 mRNA, complete cds. /FEA=mRNA /PROD=PRO0872 /DB_XREF=gi:11493406 /UG=Hs.172154 Homo sapiens clone FLB3442	AF130050	1.07
hypothetical protein FLJ10743	AA915989	1.07
microtubule-associated protein 7	AW242297	1.07
amiloride-sensitive cation channel 3, testis	NM_020322	1.07

TRK-fused gene	NM_006070	1.07
ATPase inhibitor precursor	NM_016311	1.07
vesicle-associated membrane protein 3 (cellubrevin)	BC005941	1.07
DKFZP547E2110 protein	NM_014034	1.07
insulin upstream factor 1	NM_013311	1.07
phosphatidylinositol-4-phosphate 5-kinase, type I, alpha	U78577	1.07
dihydropyrimidinase-like 2	NM_001386	1.07
secreted and transmembrane 1	BF939675	1.07
runt-related transcription factor 3	AA541630	1.07
hypothetical protein LOC55565	AF070588	1.07
chromosome 11 open reading frame 5	NM_014205	1.07
polymerase (DNA directed), theta	NM_006596	1.07
Consensus includes gb:AA927664 /FEA=EST /DB_XREF=gi:3076484 /DB_XREF=est:om71h10.s1 /CLONE=IMAGE:1552675 /UG=Hs.171626 transcription elongation factor B (SIII), polypep	AA927664	1.07
heterogeneous nuclear ribonucleoprotein A1	AI144007	1.07
hypothetical protein FLJ20136	NM_017684	1.07
bone morphogenetic protein 8 (osteogenic protein 2)	NM_001720	1.07
prefoldin 1	NM_002622	1.07
hypothetical protein FLJ10956	NM_018283	1.07
steroid-5-alpha-reductase, alpha polypeptide 1 (3-oxo-5 alpha-steroid delta 4-dehydrogenase alpha 1)	AF113128	1.07
cysteine-rich, angiogenic inducer, 61	AF003114	1.07
zinc finger protein 220	NM_006766	1.07
BAI1-associated protein 3	NM_003933	1.07
dyskeratosis congenita 1, dyskerin	AI010395	1.07
hypothetical protein FLJ20085	NM_017660	1.07
formin-like	NM_005892	1.07
Consensus includes gb:AK026803.1 /DEF=Homo sapiens cDNA: FLJ23150 fis, clone LNG09358. /FEA=mRNA /DB_XREF=gi:10439742 /UG=Hs.277176 Homo sapiens cDNA: FLJ23150 fis, cl	AK026803	1.07
glycerol kinase	AI830490	1.07
hypothetical protein FLJ20259	BC000978	1.07
N-acetylgalactosaminidase, alpha-	M38083	1.07
poly(A) binding protein, cytoplasmic 3	NM_030979	1.07
GTP-binding protein rag8	NM_016656	1.07
kinectin 1 (kinesin receptor)	Z22551	1.07
DKFZP434P1750 protein	NM_015527	1.07
RAD9 homolog (S. pombe)	NM_004584	1.07
nitric oxide synthase 2A (inducible, hepatocytes)	L24553	1.07
solute carrier family 25 (carnitine/acylcarnitine translocase), member 20	BC001689	1.07
hypothetical protein FLJ11336	NM_018393	1.07
protein inhibitor of activated STAT3	NM_006099	1.07
protein tyrosine phosphatase, non-receptor type substrate 1	D86043	1.07
fragile X mental retardation, autosomal homolog 1	BG025078	1.07
zinc finger RNA binding protein	AI459274	1.07
amyotrophic lateral sclerosis 2 (juvenile) chromosome region, candidate 3	AV705253	1.07
hypothetical protein FLJ20005	NM_017617	1.07
glioblastoma amplified sequence	NM_001483	1.07
Alport syndrome, mental retardation, midface hypoplasia and elliptocytosis chromosomal region, gene 1	AK023637	1.07
solute carrier family 25 (mitochondrial carrier, Aralar), member 12	NM_003705	1.07
COQ7 coenzyme Q, 7 homolog ubiquinone (yeast)	AF032900	1.06
nuclear receptor subfamily 4, group A, member 1	D49728	1.06
potassium voltage-gated channel, Shaw-related subfamily, member 3	NM_004977	1.06
N-deacetylase/N-sulfotransferase (heparan glucosaminyl) 1	NM_001543	1.06
gb:L21756.1 /DEF=Homo sapiens acute myeloid leukemia associated protein (AML1EAP) mRNA, complete cds. /FEA=mRNA /GEN=AML1EAP /DB_XREF=gi:400340 /FL=gb:L21756.1	L21756	1.06
guanine nucleotide binding protein (G protein), gamma 7	BF432795	1.06
KIAA0997 protein	AI656706	1.06
sin3-associated polypeptide, 30kD	AF055993	1.06
spondyloepiphyseal dysplasia, late, pseudogene	AF291676	1.06
translocase of outer mitochondrial membrane 70 homolog A (yeast)	BC003633	1.06
myo-inositol 1-phosphate synthase A1	AL137749	1.06
glucosamine (N-acetyl)-6-sulfatase (Sanfilippo disease IIID)	NM_002076	1.06
ubiquitin specific protease 4 (proto-oncogene)	AF017306	1.06
KIAA0100 gene product	AA904674	1.06
interleukin 10 receptor, alpha	NM_001558	1.06
hypothetical protein	NM_017571	1.06
hypothetical protein FLJ11722	NM_024970	1.06

TIA1 cytotoxic granule-associated RNA binding protein	H96549	1.06
hypothetical protein FLJ13593	NM_024780	1.06
endothelial cell growth factor 1 (platelet-derived)	NM_001953	1.06
arachidonate 12-lipoxygenase pseudogene 2	AF020774	1.06
RAB11A, member RAS oncogene family	NM_004663	1.06
small inducible cytokine subfamily B (Cys-X-Cys), member 11	AF030514	1.06
dimethylarginine dimethylaminohydrolase 2	AK026191	1.06
Consensus includes gb:AU146809 /FEA=EST /DB_XREF=gi:11008330 /DB_XREF=est:AU146809 /CLONE=HEMBB1001564 /UG=Hs.287476 Homo sapiens cDNA FLJ12004 fis, clone HEMBB	AU146809	1.06
Ca2+-dependent activator protein for secretion	NM_003716	1.06
cadherin 13, H-cadherin (heart)	NM_001257	1.06
tripartite motif-containing 3	AI419307	1.06
aspartoacylase (aminoacylase 2, Canavan disease)	NM_000049	1.06
translocating chain-associating membrane protein	BC000687	1.06
sorting nexin 10	NM_013322	1.06
ring finger protein 1	NM_002931	1.06
hypothetical protein FLJ23119	NM_024652	1.06
gamma-aminobutyric acid (GABA) A receptor, gamma 3	S82769	1.06
membrane-bound transcription factor protease, site 2	NM_015884	1.06
stomatin (EBP72)-like 1	NM_004809	1.06
novel protein	NM_017515	1.06
annexin A7	NM_004034	1.06
hypothetical protein FLJ23445	NM_025075	1.06
B-cell CLL/lymphoma 11A (zinc finger protein)	AF080216	1.06
dynein, axonemal, heavy polypeptide 9	AF257737	1.06
period homolog 1 (Drosophila)	AF022991	1.06
chromosome 18 open reading frame 1	AW008505	1.06
hypothetical protein similar to preferentially expressed antigen of melanoma	NM_023013	1.06
polycystic kidney disease 2-like 1	NM_016112	1.06
maternally expressed 3	AF090934	1.06
hypothetical protein DKFZp566I133	NM_030938	1.06
KIAA0802 protein	AB018345	1.06
mannan-binding lectin serine protease 1 (C4/C2 activating component of Ra-reactive factor)	D17525	1.06
phorbolin-like protein MDS019	BF508948	1.06
low density lipoprotein receptor (familial hypercholesterolemia)	S70123	1.06
myosin, light polypeptide 4, alkali; atrial, embryonic	AI133353	1.06
hypothetical protein FLJ20116	NM_017671	1.06
popeye protein 3	AI926544	1.06
Consensus includes gb:AL138761 /DEF=Human DNA sequence from clone RP11-16H23 on chromosome 10. Contains the gene KIAA0204 (HSLK) for a protein kinase, the COL17A1 gene for	AL138761	1.06
nuclear receptor coactivator 2	AI040324	1.06
thrombopoietin (myeloproliferative leukemia virus oncogene ligand, megakaryocyte growth and development factor)	U59495	1.06
nucleolar protein ANKT	NM_018454	1.06
coactivator-associated arginine methyltransferase-1	AA551784	1.06
myeloid/lymphoid or mixed-lineage leukemia (trithorax homolog, Drosophila); translocated to, 2	L22179	1.06
interferon, gamma-inducible protein 16	BG256677	1.06
peptidylprolyl isomerase (cyclophilin)-like 2	NM_014337	1.06
Consensus includes gb:AI095896 /FEA=EST /DB_XREF=gi:3434872 /DB_XREF=est:qb21a10.x1 /CLONE=IMAGE:1696890 /UG=Hs.301463 Human Chromosome 16 BAC clone CIT987SK-A-6	AI095896	1.06
RAD51-like 3 (S. cerevisiae)	AB016223	1.06
LIS1-interacting protein NUDEL; endooligopeptidase A	NM_030808	1.06
CGI-96 protein	AL590118	1.06
hypothetical protein DKFZp762I166	AI341234	1.06
G protein-coupled receptor 51	AF069755	1.06
HSPC055 protein	NM_014153	1.06
ezrin	AA196245	1.06
chaperonin containing TCP1, subunit 5 (epsilon)	AF275798	1.06
ephrin-A4	NM_005227	1.06
splicing factor proline/glutamine rich (polypyrimidine tract binding protein associated)	AV705803	1.06
ceroid-lipofuscinosis, neuronal 8 (epilepsy, progressive with mental retardation)	NM_018941	1.06
homeo box 11-like 2	NM_021025	1.06
protein kinase, cAMP-dependent, regulatory, type II, alpha	NM_004157	1.06
TP53 target gene 1	AB007457	1.06
neurexin 3	NM_004796	1.06
RAN binding protein 17	NM_022897	1.06
transducin (beta)-like 1	AW968555	1.06

nuclear receptor subfamily 0, group B, member 1	NM_000475	1.06
KIAA1115 protein	BC002799	1.06
lamin B receptor	NM_002296	1.06
ATPase, H+ transporting lysosomal (vacuolar proton pump), member M	AF077614	1.06
aspartylglucosaminidase	M64073	1.06
heat shock 70kD protein 4	AA043348	1.06
high-glucose-regulated protein 8	NM_016258	1.06
mitochondrial intermediate peptidase	U80034	1.06
dolichyl-phosphate mannosyltransferase polypeptide 1, catalytic subunit	NM_003859	1.06
Consensus includes gb:BF725121 /FEA=EST /DB_XREF=gi:12041032 /DB_XREF=est:bx12e01.x1 /CLONE=bx12e01 /UG=Hs.272108 ESTs	BF725121	1.06
Consensus includes gb:AA343027 /FEA=EST /DB_XREF=gi:1995263 /DB_XREF=est:EST48665 /UG=Hs.193606 Homo sapiens PAC clone RP5-1093017 from 7q11.23-q21	AA343027	1.06
deltex homolog 2 (Drosophila)	AK023924	1.06
hypothetical protein MGC4179	NM_024340	1.06
nephroblastoma overexpressed gene	NM_002514	1.06
vesicle-associated membrane protein 1 (synaptobrevin 1)	AU150319	1.06
carboxypeptidase D	D85390	1.06
f-box and WD-40 domain protein 1B	BE963245	1.06
dual specificity phosphatase 11 (RNA/RNP complex 1-interacting)	NM_003584	1.06
DKFZP564F0522 protein	AL049943	1.06
solute carrier family 16 (monocarboxylic acid transporters), member 5	NM_004695	1.06
Consensus includes gb:BE549732 /FEA=EST /DB_XREF=gi:9791424 /DB_XREF=est:7b38a04.x1 /CLONE=IMAGE:3230478 /UG=Hs.112158 Homo sapiens PAC clone RP4-751H13 from 7q3	BE549732	1.06
likely ortholog of mouse variant polyadenylation protein CSTF-64	AI872408	1.06
solute carrier family 9 (sodium/hydrogen exchanger), isoform 6	NM_006359	1.06
nuclear factor I/C (CCAAT-binding transcription factor)	NM_005597	1.06
carboxypeptidase M	AV710357	1.06
Consensus includes gb:BF448315 /FEA=EST /DB_XREF=gi:11513897 /DB_XREF=est:nad18a03.x1 /CLONE=IMAGE:3365668 /UG=Hs.152939 Homo sapiens clone 24630 mRNA sequence	BF448315	1.06
BCL2/adenovirus E1B 19kD interacting protein 1	NM_013979	1.06
transforming growth factor, beta 1	NM_000660	1.06
Consensus includes gb:X84340.1 /DEF=H.sapiens mRNA for Ig light chain, variable region (ID:CLL001VL). /FEA=mRNA /GEN=Ig VL gene /PROD=immunoglobulin light chain variable region .	X84340	1.06
ankylosis, progressive homolog (mouse)	NM_019847	1.06
bridging integrator 1	AF001383	1.06
Consensus includes gb:AI073983 /FEA=EST /DB_XREF=gi:3400627 /DB_XREF=est:oy66c04.x1 /CLONE=IMAGE:1670790 /UG=Hs.298553 Homo sapiens, clone IMAGE:3953631, mRNA, pa	AI073983	1.06
thrombopoietin (myeloproliferative leukemia virus oncogene ligand, megakaryocyte growth and development factor)	D32047	1.06
protein tyrosine phosphatase, non-receptor type 21	X79510	1.06
pyruvate dehydrogenase kinase, isoenzyme 1	NM_002610	1.06
mannosidase, alpha, class 1A, member 2	BE543064	1.06
annexin A2	BC001388	1.06
candidate tumor suppressor p33 ING1 homolog	AA887083	1.06
LAP (leucine-rich repeats and PDZ) and no PDZ protein	NM_018214	1.06
deoxyribonuclease I-like 2	NM_001374	1.06
similar to hypothetical protein MNCb-2386	BG230758	1.06
GDNF family receptor alpha 2	U97145	1.06
Consensus includes gb:AW166925 /FEA=EST /DB_XREF=gi:6398450 /DB_XREF=est:xg68e01.x1 /CLONE=IMAGE:2633496 /UG=Hs.289037 Homo sapiens cDNA FLJ14135 fis, clone MAMM/	AW166925	1.06
candidate tumor suppressor p33 ING1 homolog	NM_016162	1.06
Ras association (RalGDS/AF-6) domain family 1	NM_007182	1.06
HMT1 hnRNP methyltransferase-like 1 (S. cerevisiae)	NM_001535	1.06
cholinergic receptor, nicotinic, beta polypeptide 1 (muscle)	NM_000747	1.06
tubby like protein 1	NM_003322	1.06
Consensus includes gb:BG235929 /FEA=EST /DB_XREF=gi:12749776 /DB_XREF=est:naf18h02.x1 /CLONE=IMAGE:4141658 /UG=Hs.26034 ESTs	BG235929	1.06
ADP-ribosylation factor-like 1	AU151560	1.06
polymerase (RNA) II (DNA directed) polypeptide C (33kD)	AI192781	1.06
DKFZP564O123 protein	NM_014043	1.06
KIAA0907 protein	NM_014949	1.06
BTG family, member 2	NM_006763	1.06
hypothetical protein FLJ23469	NM_024710	1.06
solute carrier family 2 (facilitated glucose transporter), member 4	NM_001042	1.06
elastase 3B, pancreatic	NM_007352	1.06
RCD1 required for cell differentiation1 homolog (S. pombe)	AI567462	1.06
B1 for mucin	X83412	1.06
Consensus includes gb:AL122122.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434L098 (from clone DKFZp434L098). /FEA=mRNA /DB_XREF=gi:6102949 /UG=Hs.274559 Homo sapiens mR	AL122122	1.06
KIAA0930 protein	AB023147	1.06
KIAA0532 protein	AI052003	1.06
keratin 9 (epidermolytic palmoplantar keratoderma)	NM_000226	1.06

CD22 antigen	NM_001771	1.06
chloride channel 3	AI760629	1.06
hypothetical protein FLJ12438	NM_021933	1.06
growth arrest-specific 1	NM_002048	1.06
amyloid beta (A4) precursor-like protein 2	NM_001642	1.06
plexin B2	AL022328	1.06
ovarian cancer overexpressed 1	NM_015945	1.06
hypothetical protein PRO0971	NM_018569	1.06
protein tyrosine phosphatase, non-receptor type 13 (APO-1/CD95 (Fas)-associated phosphatase)	NM_006264	1.06
Consensus includes gb:AF070580.1 /DEF=Homo sapiens clone 24527 mRNA sequence. /FEA=mRNA /DB_XREF=gi:3387952 /UG=Hs.25422 Homo sapiens clone 24527 mRNA sequence	AF070580	1.06
isovaleryl Coenzyme A dehydrogenase	NM_002225	1.06
NPD009 protein	AF237813	1.06
DKFZP586F1918 protein	BE645501	1.06
hypothetical protein FLJ21919	NM_023015	1.06
CD97 antigen	NM_001784	1.06
Consensus includes gb:AU147084 /FEA=EST /DB_XREF=gi:11008605 /DB_XREF=est:AU147084 /CLONE=HEMBB1002371 /UG=Hs.287482 Homo sapiens cDNA FLJ12072 fis, clone HEMBB	AU147084	1.06
cerebroside (3'-phosphoadenylylsulfate:galactosylceramide 3') sulfotransferase	NM_004861	1.06
hypothetical protein DKFZp7628226	AK001821	1.06
HIV-1 inducer of short transcripts binding protein; lymphoma related factor	AI568395	1.06
hypothetical protein FLJ10060	NM_017986	1.06
acyl-Coenzyme A oxidase 1, palmitoyl	S69189	1.06
Consensus includes gb:AW972351 /FEA=EST /DB_XREF=gi:8162197 /DB_XREF=est:EST384442 /UG=Hs.293451 ESTs	AW972351	1.06
ubiquitin specific protease 25	NM_013396	1.06
KIAA0397 gene product	AB007857	1.06
mitochondrial ribosomal protein S10	NM_018141	1.05
Consensus includes gb:AK024860.1 /DEF=Homo sapiens cDNA: FLJ21207 fis, clone COL00362. /FEA=mRNA /DB_XREF=gi:10437268 /UG=Hs.306708 Homo sapiens cDNA: FLJ21207 fis, cl	AK024860	1.05
small inducible cytokine subfamily A (Cys-Cys), member 25	NM_005624	1.05
KIAA0564 protein	AI744626	1.05
hypothetical protein FLJ11806	NM_024824	1.05
chromodomain helicase DNA binding protein 4	AI613273	1.05
ATP binding protein associated with cell differentiation	NM_005783	1.05
hypothetical protein MGC2963	NM_031298	1.05
KIAA0323 protein	AB002321	1.05
hypothetical protein MGC3262	NM_024029	1.05
hypothetical protein MGC2744	AI093187	1.05
myotubularin related protein 1	AF057354	1.05
RecQ protein-like (DNA helicase Q1-like)	AI685944	1.05
Consensus includes gb:AK022331.1 /DEF=Homo sapiens cDNA FLJ12269 fis, clone MAMMA1001635. /FEA=mRNA /DB_XREF=gi:10433706 /UG=Hs.293925 Homo sapiens cDNA FLJ12269 f	AK022331	1.05
interferon (alpha, beta and omega) receptor 1	NM_000629	1.05
for protein disulfide isomerase-related	NM_006810	1.05
KIAA1065 protein	NM_014964	1.05
hypothetical protein FLJ13920	NM_024558	1.05
tumor differentially expressed 1	AW173623	1.05
hypothetical protein	AI800419	1.05
testis derived transcript (3 LIM domains)	BC001451	1.05
zinc finger protein 226	NM_015919	1.05
hypothetical protein	AI928407	1.05
PFTAIRE protein kinase 1	NM_012395	1.05
UDP glycosyltransferase 1 family, polypeptide A9	NM_021027	1.05
POU domain, class 3, transcription factor 2	NM_005604	1.05
ISL1 transcription factor, LIM/homeodomain, (islet-1)	NM_002202	1.05
biotinidase	AI767414	1.05
methylcrotonoyl-Coenzyme A carboxylase 1 (alpha)	NM_020166	1.05
hypothetical protein DKFZp434E1723	AI743331	1.05
trinucleotide repeat containing 15	AL045800	1.05
docking protein 1, 62kD (downstream of tyrosine kinase 1)	AF180527	1.05
S100 calcium binding protein A1	NM_006271	1.05
synaptosomal-associated protein, 29kD	NM_004782	1.05
cullin 3	AF062537	1.05
CASP8 associated protein 2	AB037736	1.05
KIAA0013 gene product	NM_014783	1.05
nuclear factor I/B	U70862	1.05
hypothetical protein FLJ12565	NM_022064	1.05

hypothetical protein FLJ22625	NM_024715	1.05
TATA box binding protein (TBP)-associated factor, RNA polymerase II, S, 150kD	NM_006706	1.05
putative translation initiation factor	W28892	1.05
TRAF interacting protein	NM_005879	1.05
myotubularin related protein 2	AK027038	1.05
huntingtin (Huntington disease)	NM_002111	1.05
stathmin-like 3	NM_015894	1.05
E2F transcription factor 6	NM_001952	1.05
hypothetical protein FLJ22059	AV756161	1.05
lymphocyte antigen 117	AV713720	1.05
serine protease inhibitor, Kunitz type 1	NM_003710	1.05
CDP-diacylglycerol synthase (phosphatidate cytidylyltransferase) 2	Y16521	1.05
Consensus includes gb:AL109706.1 /DEF=Homo sapiens mRNA full length insert cDNA clone EUROIMAGE 362430. /FEA=mRNA /DB_XREF=gi:5689823 /UG=Hs.32769 Homo sapiens mRNA	AL109706	1.05
KIAA1054 protein	AB028977	1.05
collagen, type II, alpha 1 (primary osteoarthritis, spondyloepiphyseal dysplasia, congenital)	X06268	1.05
hypothetical protein FLJ12568	NM_024993	1.05
collectin sub-family member 10 (C-type lectin)	NM_006438	1.05
translin-associated factor X	NM_005999	1.05
RAB32, member RAS oncogene family	NM_006834	1.05
NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 5 (13kD, B13)	AK022209	1.05
dystrobrevin, alpha	U46746	1.05
gb:NM_017871.1 /DEF=Homo sapiens hypothetical protein FLJ20542 (FLJ20542), mRNA. /FEA=mRNA /GEN=FLJ20542 /PROD=hypothetical protein FLJ20542 /DB_XREF=gi:8923511 /UG=	NM_017871	1.05
XIAP associated factor-1	NM_017523	1.05
KIAA0586 gene product	NM_014749	1.05
pyruvate kinase, liver and RBC	NM_000298	1.05
ADP-ribosylation factor 4	NM_001660	1.05
Consensus includes gb:AL049268.1 /DEF=Homo sapiens mRNA; cDNA DKFZp564G103 (from clone DKFZp564G103). /FEA=mRNA /DB_XREF=gi:4500017 /UG=Hs.202684 Homo sapiens mI	AL049268	1.05
calcyon; D1 dopamine receptor-interacting protein	NM_015722	1.05
synaptic vesicle protein 2B homolog	NM_014848	1.05
FK506 binding protein 8 (38kD)	L37033	1.05
CD6 antigen	AW134823	1.05
KIAA0266 gene product	NM_021645	1.05
BRF2, subunit of RNA polymerase III transcription initiation factor, BRF1-like	NM_018310	1.05
catenin, beta interacting protein 1	NM_020248	1.05
KIAA0759 protein	AB018302	1.05
xylulokinase (H. influenzae) homolog	NM_005108	1.05
nuclear factor (erythroid-derived 2)-like 1	NM_003204	1.05
fetal hypothetical protein	NM_013303	1.05
Consensus includes gb:AW269792 /FEA=EST /DB_XREF=gi:6656822 /DB_XREF=est:xv45d03.x1 /CLONE=IMAGE:2816069 /UG=Hs.5010 Homo sapiens clone 24672 mRNA sequence	AW269792	1.05
cytochrome P450, subfamily IIA (phenobarbital-inducible), polypeptide 7, pseudogene 2	U22044	1.05
hypothetical protein FLJ13189	NM_024682	1.05
vitamin D (1,25- dihydroxyvitamin D3) receptor	AA772285	1.05
mitochondrial GTP binding protein	AL524262	1.05
nesca protein	NM_014328	1.05
DKFZP434H132 protein	AW516789	1.05
antigen p97 (melanoma associated) identified by monoclonal antibodies 133.2 and 96.5	NM_005929	1.05
Kelch-like ECH-associated protein 1	NM_012289	1.05
2',3'-cyclic nucleotide 3' phosphodiesterase	BC001362	1.05
hypothetical protein FLJ10120	AI140364	1.05
bromodomain adjacent to zinc finger domain, 2B	NM_013450	1.05
CGI-96 protein	Z93241	1.05
RAS p21 protein activator (GTPase activating protein) 3 (Ins(1,3,4,5)P4-binding protein)	NM_007368	1.05
tRNA selenocysteine associated protein	NM_017846	1.05
hypothetical protein FLJ10157	NM_018012	1.05
gastric inhibitory polypeptide	NM_004123	1.05
caspase 7, apoptosis-related cysteine protease	NM_001227	1.05
potassium voltage-gated channel, subfamily H (eag-related), member 2	AB044806	1.05
p10-binding protein	NM_024491	1.05
lipin 1	AV705559	1.05
excision repair cross-complementing rodent repair deficiency, complementation group 2 (xeroderma pigmentosum D)	AI918117	1.05
CASP8 and FADD-like apoptosis regulator	AF041459	1.05
PMS1 postmeiotic segregation increased 1 (S. cerevisiae)	BG434893	1.05
hypothetical protein, estradiol-induced	NM_015516	1.05

Consensus includes gb:AL031178 /DEF=Human DNA sequence from clone 341E18 on chromosome 6p11.2-12.3. Contains a SerineThreonine Protein Kinase gene (presumptive isolog of a R	AL031178	1.05
retinoblastoma binding protein 2	NM_005056	1.05
interferon gamma receptor 2 (interferon gamma transducer 1)	NM_005534	1.05
Consensus includes gb:AF007131.1 /DEF=Homo sapiens clone 23963 mRNA sequence. /FEA=mRNA /DB_XREF=gi:2852605 /UG=Hs.48483 Homo sapiens clone 23963 mRNA sequence	AF007131	1.05
sorting nexin 2	NM_003100	1.05
MADS box transcription enhancer factor 2, polypeptide A (myocyte enhancer factor 2A)	NM_005587	1.05
hypothetical protein FLJ10143	NM_018009	1.05
hemochromatosis	AF144241	1.05
Consensus includes gb:AW519168 /FEA=EST /DB_XREF=gi:7157250 /DB_XREF=est:ha52a08.x1 /CLONE=IMAGE:2877302 /UG=Hs.163271 Human alpha-1 Ig germline C-region membrane-	AW519168	1.05
solute carrier family 6 (neurotransmitter transporter, taurine), member 6	NM_003043	1.05
membrane protein of cholinergic synaptic vesicles	BC001913	1.05
solute carrier family 5 (sodium iodide symporter), member 5	D87920	1.05
zinc finger protein 263	NM_005741	1.05
protein tyrosine phosphatase, non-receptor type 4 (megakaryocyte)	NM_002830	1.05
E2F transcription factor 3	NM_001949	1.05
guanine nucleotide binding protein (G protein), q polypeptide	U40038	1.05
neurotrophin 3	NM_002527	1.05
15 kDa selenoprotein	NM_004261	1.05
hypothetical protein FLJ12517	NM_023007	1.05
protein O-fucosyltransferase 1	BC000582	1.05
sec22 homolog	NM_012430	1.05
Consensus includes gb:AL049387.1 /DEF=Homo sapiens mRNA; cDNA DKFZp586N1918 (from clone DKFZp586N1918). /FEA=mRNA /DB_XREF=gi:4500178 /UG=Hs.169936 Homo sapiens	AL049387	1.05
protein phosphatase 1, catalytic subunit, gamma isoform	NM_002710	1.05
hypothetical protein FLJ12488	NM_031218	1.05
cathepsin S	NM_004079	1.05
death effector domain-containing	AK022531	1.05
potassium voltage-gated channel, KQT-like subfamily, member 1	NM_000218	1.05
leukocyte receptor cluster (LRC) member 5	NM_024075	1.05
immunoglobulin lambda locus	X79782	1.05
protein phosphatase 4 regulatory subunit 2	AI582773	1.05
ATP-binding cassette, sub-family F (GCN20), member 2	NM_005692	1.05
beta-transducin repeat containing	AF101784	1.05
hypothetical protein MGC5508	NM_024092	1.05
erythroid differentiation and denucleation factor 1	NM_006553	1.05
RABSC, member RAS oncogene family	AF141304	1.05
KIAA0542 gene product	AB011114	1.05
hypothetical protein FLJ20320	NM_017765	1.05
hypothetical protein MGC4645	NM_024515	1.05
BN51 (BHK21) temperature sensitivity complementing	NM_001722	1.05
multiple endocrine neoplasia I	NM_000244	1.05
hypothetical protein FLJ11856	NM_024531	1.05
NAD(P)H dehydrogenase, quinone 1	BC000906	1.05
solute carrier family 11 (proton-coupled divalent metal ion transporters), member 2	AF046997	1.05
hypothetical protein FLJ23132	NM_025054	1.05
sarcosine dehydrogenase	AF047190	1.05
ubiquitin-conjugating enzyme E2H (UBC8 homolog, yeast)	NM_003344	1.05
breast carcinoma amplified sequence 2	NM_005872	1.05
ras responsive element binding protein 1	AF072826	1.05
Fc fragment of IgA, receptor for	U43677	1.05
Consensus includes gb:AP000693 /DEF=Homo sapiens genomic DNA, chromosome 21q22.2, PCR fragment from BAC clone:KB739C11, CBR1-HLCS region /FEA=mRNA /DB_XREF=gi:669363	AP000693	1.05
Consensus includes gb:AI928387 /FEA=EST /DB_XREF=gi:5664351 /DB_XREF=est:wo96f10.x1 /CLONE=IMAGE:2463211 /UG=Hs.331601 Homo sapiens, Similar to cysteine and histidine-r	AI928387	1.05
F-box only protein 4	NM_018007	1.05
KIAA0133 gene product	NM_014777	1.05
UBX domain-containing 2	AI927512	1.05
mitochondrial ribosomal protein S31	NM_005830	1.05
tumor necrosis factor (ligand) superfamily, member 6	AF288573	1.04
6-pyruvoyl-tetrahydropterin synthase/dimerization cofactor of hepatocyte nuclear factor 1 alpha (TCF1)	NM_000281	1.04
guanylate cyclase 1, soluble, alpha 2	NM_000855	1.04
signal transducer and activator of transcription 5B	NM_012448	1.04
hypothetical protein FLJ20113	NM_017670	1.04
galactokinase 1	BG474736	1.04
cytoplasmic FMRP interacting protein 2	AL161999	1.04
Consensus includes gb:AL109714.1 /DEF=Homo sapiens mRNA full length insert cDNA clone EUROIMAGE 327506. /FEA=mRNA /DB_XREF=gi:5689826 /UG=Hs.195484 Homo sapiens mRN	AL109714	1.04

popeye protein 3	AV703259	1.04
kinase insert domain receptor (a type III receptor tyrosine kinase)	NM_002253	1.04
exostoses (multiple) 2	NM_000401	1.04
interferon, gamma-inducible protein 16	NM_005531	1.04
hypothetical protein FLJ22195	AL523965	1.04
Consensus includes gb:Y08772 /DEF=H.sapiens PABII pseudogene /FEA=CDS /DB_XREF=gi:1770485 /UG=Hs.247983 H.sapiens PABII pseudogene	Y08772	1.04
a disintegrin and metalloproteinase domain 19 (meltrin beta)	Y13786	1.04
similar to granzyme B (granzyme 2, cytotoxic T-lymphocyte-associated serine esterase 1) (H. sapiens)	M36118	1.04
CUG triplet repeat, RNA binding protein 1	N25915	1.04
KIAA0254 gene product	NM_014758	1.04
heat shock 70kD protein 4	AB023420	1.04
Consensus includes gb:AK022806.1 /DEF=Homo sapiens cDNA FLJ12744 fis, clone NT2RP2000715. /FEA=mRNA /DB_XREF=gi:10434418 /UG=Hs.139425 Homo sapiens cDNA FLJ12744 fi	AK022806	1.04
FGF receptor activating protein 1	AL049261	1.04
v-ski sarcoma viral oncogene homolog (avian)	AI568728	1.04
fucosidase, alpha-L- 1, tissue	NM_000147	1.04
protease, serine, 2 (trypsin 2)	AF009664	1.04
glycosylphosphatidylinositol specific phospholipase D1	AL031230	1.04
2,3-bisphosphoglycerate mutase	NM_001724	1.04
KIAA0737 gene product	BF001668	1.04
Consensus includes gb:S76346.1 /DEF=AML1=AML1 (alternatively spliced, exons 5 and b) human, mRNA Partial, 284 nt. /FEA=mRNA /GEN=AML1 /PROD=AML1 /DB_XREF=gi:913288 /UG	S76346	1.04
synovial sarcoma, X breakpoint 1	BC001003	1.04
hypothetical protein FLJ22237	NM_024827	1.04
KIAA0652 gene product	BC001331	1.04
lymphoid enhancer-binding factor 1	AF288571	1.04
phosphodiesterase 4D interacting protein (myomegalin)	AB007923	1.04
proenkephalin	NM_006211	1.04
collagen, type IV, alpha 6	AL031177	1.04
nucleosomal binding protein 1	BC005342	1.04
acetyl-Coenzyme A carboxylase alpha	BE855983	1.04
transcription factor AP-2 gamma (activating enhancer binding protein 2 gamma)	U85658	1.04
interleukin 8 receptor, alpha	NM_000634	1.04
hypothetical protein FLJ23323	NM_024654	1.04
E3 ubiquitin ligase SMURF1	AI654869	1.04
STRIN protein	NM_016271	1.04
Consensus includes gb:AL359931.1 /DEF=Homo sapiens mRNA full length insert cDNA clone EUROIMAGE 288936. /FEA=mRNA /DB_XREF=gi:8977886 /UG=Hs.48607 Homo sapiens mRNA	AL359931	1.04
RNA cyclase homolog	NM_005772	1.04
papillomavirus regulatory factor PRF-1	NM_018660	1.04
KIAA1096 protein	AL096857	1.04
hypothetical protein FLJ12799	NM_022495	1.04
hypothetical protein FLJ10700	NM_018182	1.04
retinoschisis (X-linked, juvenile) 1	AL049684	1.04
hypothetical protein PRO2964	NM_018547	1.04
interleukin 1 receptor-like 1	AB012701	1.04
KIAA1735 protein	AF070621	1.04
complement C1r-like proteinase precursor,	NM_016546	1.04
glucocorticoid modulatory element binding protein 2	AA045183	1.04
RNA helicase-related protein	AF078844	1.04
ets variant gene 2	AV693985	1.04
hypothetical protein FLJ11137	NM_018337	1.04
smoothelin	AF064238	1.04
hypothetical protein FLJ11323	NM_018390	1.04
KIAA0335 gene product	NM_014803	1.04
discs, large (Drosophila) homolog-associated protein 1	AB000277	1.04
hypothetical protein	NM_015684	1.04
Consensus includes gb:AI350500 /FEA=EST /DB_XREF=gi:4087706 /DB_XREF=est:qt17h03.x1 /CLONE=IMAGE:1947893 /UG=Hs.301463 Human Chromosome 16 BAC clone CIT987SK-A-6	AI350500	1.04
sorting nexin 11	AI668643	1.04
KIAA0349 protein	AI761518	1.04
ilvB (bacterial acetolactate synthase)-like	AC004794	1.04
hypothetical protein FLJ11467	AL137434	1.04
ubiquitin-conjugating enzyme E2H (UBC8 homolog, yeast)	AI829920	1.04
ATP-binding cassette, sub-family F (GCN20), member 2	BC001661	1.04
2'-5'-oligoadenylate synthetase 3 (100 kD)	NM_006187	1.04
methyl CpG binding protein 2 (Rett syndrome)	L37298	1.04

Consensus includes gb:AK025343.1 /DEF=Homo sapiens cDNA: FLJ21690 fis, clone COL09538. /FEA=mRNA /DB_XREF=gi:10437841 /UG=Hs.306802 Homo sapiens cDNA: FLJ21690 fis, cl	AK025343	1.04
splicing factor, arginine/serine-rich 7 (35kD)	NM_006276	1.04
Consensus includes gb:AW274747 /FEA=EST /DB_XREF=gi:6661777 /DB_XREF=est:xn34e08.x1 /CLONE=IMAGE:2695622 /UG=Hs.270527 Homo sapiens mRNA; cDNA DKFZp434F0327 (fr	AW274747	1.04
calcium channel, voltage-dependent, alpha 1G subunit	AB012043	1.04
hypothetical protein FLJ13154	NM_024598	1.04
vesicle-associated membrane protein 5 (myobrevin)	NM_006634	1.04
HSPC034 protein	AV700696	1.04
Consensus includes gb:AF229166.1 /DEF=Homo sapiens docking protein 1-like protein mRNA, partial cds. /FEA=mRNA /PROD=docking protein 1-like protein /DB_XREF=gi:7650361 /UG=I	AF229166	1.04
eukaryotic translation initiation factor 2-alpha kinase 3	NM_004836	1.04
KIAA0923 protein	AL133046	1.04
6-phosphofructo-2-kinase/fructose-2,6-biphosphatase 2	NM_006212	1.04
KIAA0418 gene product	NM_014631	1.04
LPS-induced TNF-alpha factor	AB034747	1.04
hypothetical protein FLJ20719	AV684285	1.04
KIAA0280 protein	D87470	1.04
C6l-147 protein	NM_016077	1.04
Consensus includes gb:BG500611 /FEA=EST /DB_XREF=gi:13462128 /DB_XREF=est:602546082F1 /CLONE=IMAGE:4668261 /UG=Hs.82719 Homo sapiens mRNA; cDNA DKFZp586F1822	BG500611	1.04
carbonic anhydrase VB, mitochondrial	NM_007220	1.04
phosphodiesterase 2A, cGMP-stimulated	NM_002599	1.04
hypothetical protein MGC9084	AL035369	1.04
hypothetical protein FLJ20039	NM_017635	1.04
calcium/calmodulin-dependent protein kinase (CaM kinase) II beta	AF112472	1.04
hypothetical protein FLJ10213	NM_018029	1.04
zinc finger protein 75 (D8C6)	W90796	1.04
KIAA1882 protein	AW291218	1.04
ADP-ribosylation factor GTPase activating protein 1	BC005122	1.04
KIAA0226 gene product	BF448041	1.04
DAZ associated protein 2	N34846	1.04
mitogen-activated protein kinase kinase 5	NM_002757	1.04
calpain 10	NM_023089	1.04
ATP-binding cassette, sub-family C (CFTR/MRP), member 5	AF146074	1.04
ribosomal protein L27	NM_000988	1.04
Toll-interacting protein	NM_019009	1.04
Consensus includes gb:AU157915 /FEA=EST /DB_XREF=gi:11019436 /DB_XREF=est:AU157915 /CLONE=PLACE1010102 /UG=Hs.5518 Homo sapiens cDNA FLJ11311 fis, clone PLACE101(	AU157915	1.04
homeo box C4	AW299598	1.04
KIAA0616 protein	AB014516	1.04
KIAA0240 protein	D87077	1.04
prostaglandin E receptor 4 (subtype EP4)	AI675173	1.04
oxygen regulated protein (150kD)	NM_006389	1.04
colony stimulating factor 2 (granulocyte-macrophage)	M11734	1.04
nuclear receptor subfamily 2, group E, member 3	NM_014249	1.04
tumor protein 63 kDa with strong homology to p53	AB010153	1.04
transmembrane 4 superfamily member 5	NM_003963	1.04
Cluster Incl. AA527515:ng41h06.s1 Homo sapiens cDNA, 3' end /clone=IMAGE-937403 /clone_end=3' /gb=AA527515 /gi=2269584 /ug=Hs.238570 /len=587	AA527515	1.04
DIPB protein	BF431488	1.04
Consensus includes gb:AI249275 /DEF=Homo sapiens partial MTHFR gene for methylenetetrahydrofolate reductase /FEA=CDS /DB_XREF=gi:7271187 /UG=Hs.214142 5,10-methylenetetra	AI249275	1.04
SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily a, member 5	AI652586	1.04
solute carrier family 7 (cationic amino acid transporter, y+ system), member 6	NM_003983	1.04
fucosyltransferase 2 (secretor status included)	BC001899	1.04
splicing factor, arginine/serine-rich 10 (transformer 2 homolog, Drosophila)	NM_004593	1.04
BET1 homolog (S. cerevisiae)	BC000899	1.04
gb:AY027523.1 /DEF=Homo sapiens transcription factor TZP (TZP) mRNA, complete cds. /FEA=mRNA /GEN=TZP /PROD=transcription factor TZP /DB_XREF=gi:13195150 /UG=Hs.112594	AY027523	1.04
embryonic ectoderm development	AF099032	1.04
KIAA0768 protein	AF307080	1.04
CD59 antigen p18-20 (antigen identified by monoclonal antibodies 16.3A5, EJ16, EJ30, EL32 and G344)	X16447	1.04
RAB2, member RAS oncogene family	AA535244	1.04
hypothetical protein FLJ10815	AI963454	1.04
ariadne homolog, ubiquitin-conjugating enzyme E2 binding protein, 1 (Drosophila)	AL040708	1.04
cytochrome P450, subfamily XIB (steroid 11-beta-hydroxylase), polypeptide 1	AV702430	1.04
protein tyrosine phosphatase, receptor type, J	D37781	1.04
carbohydrate (chondroitin) synthase 1	NM_014918	1.04
diacylglycerol kinase, theta (110kD)	NM_001347	1.04
coagulation factor III (thromboplastin, tissue factor)	NM_001993	1.04

tumor suppressing subtransferable candidate 1	NM_003310	1.04
f-box and leucine-rich repeat protein 5	AF157323	1.04
potassium voltage-gated channel, Isk-related family, member 1-like	NM_012282	1.04
Opa-interacting protein 5	BE045993	1.04
mannosidase, alpha, class 1A, member 1	BG287153	1.04
hypothetical protein FLJ11729	NM_024960	1.04
serine (or cysteine) proteinase inhibitor, clade A (alpha-1 antiproteinase, antitrypsin), member 2	NM_006220	1.04
odorant-binding protein 2A	NM_014582	1.04
chromosome 22 open reading frame 4	AL096779	1.04
sine oculis homeobox homolog 3 (Drosophila)	NM_005413	1.04
ADP-ribosylation factor-like 4	NM_005738	1.04
KIAA1332 protein	AK025059	1.04
a disintegrin and metalloproteinase domain 28	NM_021777	1.04
ATP-binding cassette, sub-family A (ABC1), member 2	AL162060	1.04
solute carrier family 25, member 13 (citrin)	NM_014251	1.04
zinc finger protein 36, C3H type-like 1	BE620915	1.04
solute carrier family 5 (sodium-dependent vitamin transporter), member 6	NM_021095	1.04
Consensus includes gb:AL133519 /DEF=Human DNA sequence from clone RP11-122O1 on chromosome 20 Contains a novel gene encoding a protein orthologous to the mouse vesicular inh	AL133519	1.04
cyclin D2	AW026491	1.04
collagen, type IV, alpha 2	X05610	1.04
H326	NM_015726	1.04
endosulfine alpha	AF157510	1.04
Consensus includes gb:AA114166 /FEA=EST /DB_XREF=gi:1668052 /DB_XREF=est:zn75c03.s1 /CLONE=IMAGE:564004 /UG=Hs.23964 sin3-associated polypeptide, 18kD	AA114166	1.04
tumor necrosis factor receptor superfamily, member 12 (translocating chain-association membrane protein)	U94510	1.04
polymerase (RNA) mitochondrial (DNA directed)	NM_005035	1.04
cut-like 1, CCAAT displacement protein (Drosophila)	BE046521	1.04
gb:AF132204.1 /DEF=Homo sapiens PRO2259 mRNA, complete cds. /FEA=mRNA /PROD=PRO2259 /DB_XREF=gi:11493553 /UG=Hs.302159 Homo sapiens PRO2259 mRNA, complete cds	AF132204	1.04
cAMP responsive element modulator	D14826	1.04
melanoma differentiation associated protein-5	NM_022168	1.04
nuclear receptor subfamily 1, group D, member 1	NM_021724	1.04
hypothetical protein, estradiol-induced	NM_014367	1.04
mitogen-activated protein kinase kinase kinase kinase 4	NM_004834	1.04
hypothetical protein DKFZp434L0718	AL136784	1.04
ribosomal protein S14	M13934	1.04
transcription factor 1, hepatic; LF-B1, hepatic nuclear factor (HNF1), albumin proximal factor	X71347	1.04
small nuclear RNA activating complex, polypeptide 5, 19kD	AI633709	1.04
TAF4A RNA polymerase II, TATA box binding protein (TBP)-associated factor, 135 kD	NM_003185	1.04
hypothetical protein FLJ20170	NM_017696	1.04
zinc finger protein 134 (clone pHZ-15)	AC003682	1.04
golgi-specific brefeldin A resistance factor 1	NM_004193	1.03
mitogen-activated protein kinase 8 interacting protein 1	BG222394	1.03
TdT binding protein	AL050348	1.03
ATPase, Class II, type 9B	AW411030	1.03
Consensus includes gb:AW299700 /FEA=EST /DB_XREF=gi:6709377 /DB_XREF=est:zs42g04.x1 /CLONE=IMAGE:2772342 /UG=Hs.18653 ESTs	AW299700	1.03
orosomucoid 1	NM_000607	1.03
dynein, axonemal, heavy polypeptide 9	NM_004662	1.03
KIAA0323 protein	AI075450	1.03
CGI-40 protein	NM_015996	1.03
Consensus includes gb:BE328312 /FEA=EST /DB_XREF=gi:9202173 /DB_XREF=est:hs95d12.x1 /CLONE=IMAGE:3144983 /UG=Hs.6614 ESTs, Weakly similar to A43932 mucin 2 precursor,	BE328312	1.03
secreted protein of unknown function	AA156251	1.03
KIAA0379 protein	AI081194	1.03
ribosomal protein S6 kinase, 90kD, polypeptide 3	AA906056	1.03
hypothetical protein FLJ20224	NM_017719	1.03
Consensus includes gb:AF005082.1 /DEF=Homo sapiens skin-specific protein (xp33) mRNA, partial cds. /FEA=mRNA /GEN=xp33 /PROD=skin-specific protein /DB_XREF=gi:2589191 /UG=	AF005082	1.03
mature T-cell proliferation 1	BC002600	1.03
coagulation factor VIII-associated (intronic transcript)	AJ218431	1.03
KIAA0717 protein	AY009093	1.03
solute carrier family 2 (facilitated glucose transporter), member 1	AI091047	1.03
sodium bicarbonate transporter 4	NM_021196	1.03
hematopoietic PBX-interacting protein	AI935162	1.03
protein kinase C, eta	NM_006255	1.03
nuclear receptor co-repressor 2	AI499095	1.03
angiotensin receptor 1	NM_000685	1.03

centrin, EF-hand protein, 3 (CDC31 homolog, yeast)	BC005383	1.03
hypothetical protein FLJ22060	NM_024612	1.03
protocadherin beta 6	NM_018939	1.03
paired box gene 2	AW769732	1.03
LIM domain binding 1	AF064491	1.03
Consensus includes gb:AI357616 /FEA=EST /DB_XREF=gi:4109237 /DB_XREF=est:qu20g08.s1 /CLONE=IMAGE:1965374 /UG=Hs.101651 Homo sapiens mRNA; cDNA DKFZp434C107 (fror	AI357616	1.03
hypothetical protein MGC3178	NM_030810	1.03
mitogen-activated protein kinase-activated protein kinase 5	NM_003668	1.03
brain link protein-1	NM_021817	1.03
membrane component, chromosome 17, surface marker 2 (ovarian carcinoma antigen CA125)	NM_005899	1.03
ATP-dependent RNA helicase	AF077033	1.03
Consensus includes gb:AW451236 /FEA=EST /DB_XREF=gi:6992012 /DB_XREF=est:UI-H-BI3-alh-a-07-0-UI.s1 /CLONE=IMAGE:2736661 /UG=Hs.15535 Homo sapiens clone 24582 mRNA :	AW451236	1.03
KIAA0576 protein	AB011148	1.03
ZY6 homolog	NM_006336	1.03
mitochondrial ribosomal protein L13	NM_014078	1.03
tudor repeat associator with PCTAIRE 2	AW129593	1.03
KIAA1254 protein	AB033080	1.03
hypothetical protein MGC5487	NM_024090	1.03
estrogen regulated gene 1	NM_022034	1.03
transcription factor 8 (represses interleukin 2 expression)	NM_030751	1.03
selenoprotein W, 1	NM_003009	1.03
cytochrome b-561	AL514271	1.03
advillin	AI796813	1.03
Ste-20 related kinase	NM_013233	1.03
proteasome (prosome, macropain) inhibitor subunit 1 (PI31)	NM_006814	1.03
acetylcholinesterase (YT blood group)	AI190022	1.03
ATPase, Ca++ transporting, ubiquitous	AF068220	1.03
KIAA1086 protein	AB029009	1.03
Consensus includes gb:AL049325.1 /DEF=Homo sapiens mRNA; cDNA DKFZp564D036 (from clone DKFZp564D036). /FEA=mRNA /DB_XREF=gi:4500098 /UG=Hs.112493 Homo sapiens mI	AL049325	1.03
tumor necrosis factor receptor superfamily, member 5	X60592	1.03
SMC1 structural maintenance of chromosomes 1-like 1 (yeast)	D80000	1.03
ribophorin I	NM_002950	1.03
ornithine decarboxylase antizyme 2	NM_002537	1.03
annexin A2 pseudogene 2	M62898	1.03
catenin (cadherin-associated protein), delta 1	AB002382	1.03
basic leucine zipper nuclear factor 1 (JEM-1)	NM_003666	1.03
tumor protein p53 binding protein, 2	NM_005426	1.03
ring finger protein 5	BC004155	1.03
hypothetical protein FLJ12581	NM_024865	1.03
Consensus includes gb:R37104 /FEA=EST /DB_XREF=gi:794560 /DB_XREF=est:yr53e12.s1 /CLONE=IMAGE:25731 /UG=Hs.13417 Homo sapiens clone 24670 mRNA sequence	R37104	1.03
enigma (LIM domain protein)	AI825846	1.03
cytochrome b-5	M22865	1.03
hypothetical protein	NM_019027	1.03
upstream binding protein 1 (LBP-1a)	NM_014517	1.03
hypothetical protein FLJ10661	NM_018172	1.03
solute carrier family 7 (cationic amino acid transporter, y+ system), member 8	NM_012244	1.03
nuclear receptor subfamily 2, group F, member 6	BF000629	1.03
hypothetical protein FLJ10246	NM_018038	1.03
mitogen-activated protein kinase kinase 5	U71088	1.03
calcium channel, voltage-dependent, P/Q type, alpha 1A subunit	AF004884	1.03
hypothetical protein FLJ14547	AV713053	1.03
Huntingtin interacting protein B	AI761110	1.03
hypothetical protein FLJ13852	NM_023078	1.03
solute carrier family 9 (sodium/hydrogen exchanger), isoform 3 regulatory factor 1	NM_004252	1.03
synovial sarcoma, X breakpoint 3	NM_021014	1.03
KIAA0218 gene product	NM_014760	1.03
hypothetical protein FLJ10385	AI858000	1.03
MAD, mothers against decapentaplegic homolog 2 (Drosophila)	AW151617	1.03
Consensus includes gb:AL049307.1 /DEF=Homo sapiens mRNA; cDNA DKFZp564A216 (from clone DKFZp564A216). /FEA=mRNA /DB_XREF=gi:4500076 /UG=Hs.274505 Homo sapiens mI	AL049307	1.03
hypothetical protein FLJ13490	NM_024942	1.03
phosphatidylinositol (4,5) bisphosphate 5-phosphatase homolog; phosphatidylinositol polyphosphate 5-phosphatase type IV	NM_019892	1.03
hypothetical protein FLJ10521	BC003073	1.03
hypothetical protein FLJ21128	NM_025083	1.03

CGI-10 protein	NM_015940	1.03
mitogen-activated protein kinase kinase 1	AI571419	1.03
aldehyde dehydrogenase 2 family (mitochondrial)	NM_000690	1.03
dopamine beta-hydroxylase (dopamine beta-monooxygenase)	NM_000787	1.03
KIAA0876 protein	AB020683	1.03
Burkitt lymphoma receptor 1, GTP binding protein	X68829	1.03
carcinoembryonic antigen-related cell adhesion molecule 8	M33326	1.03
hypothetical protein FLJ22457	NM_024901	1.03
hypothetical protein FLJ20424	NM_017815	1.03
single-stranded DNA-binding protein 1	BE220360	1.03
disrupted in schizophrenia 1	AK025293	1.03
fatty acid desaturase 1	BE540552	1.03
hypothetical protein FLJ22439	BF058311	1.03
cholinergic receptor, nicotinic, epsilon polypeptide	NM_000080	1.03
erythropoietin	AF053356	1.03
vascular endothelial growth factor	AF091352	1.03
protein tyrosine phosphatase, non-receptor type 21	BF116199	1.03
B-cell linker	NM_013314	1.03
chromosome 17 open reading frame 31	AB018275	1.03
zinc finger protein 36, C3H type-like 1	BG482928	1.03
bone morphogenetic protein 6	NM_001718	1.03
dUTP pyrophosphatase	U62891	1.03
MHC class II transactivator	NM_000246	1.03
paired box gene 2	NM_003988	1.03
myeloid leukemia factor 1	AI911434	1.03
KIAA0440 protein	AB007900	1.03
hypothetical protein FLJ13153	AF267859	1.03
interleukin 22 receptor	NM_021258	1.03
epiplakin 1	NM_031308	1.03
Consensus includes gb:AW167793 /FEA=EST /DB_XREF=gi:6399401 /DB_XREF=est:xg56d07.x1 /CLONE=IMAGE:2632333 /UG=Hs.164036 Homo sapiens AKAP350C mRNA sequence, alten	AW167793	1.03
KIAA0265 protein	D87454	1.03
T cell receptor delta locus	X72501	1.03
KIAA0916 protein	NM_015057	1.03
KIAA1564 protein	U00955	1.03
phosphoinositide-3-kinase, catalytic, gamma polypeptide	AF327656	1.03
heat shock 90kD protein 1, alpha	BG420237	1.03
Consensus includes gb:BF677404 /FEA=EST /DB_XREF=gi:11951299 /DB_XREF=est:602087150F1 /CLONE=IMAGE:4251205 /UG=Hs.296356 Homo sapiens mRNA; cDNA DKFZp434M162	BF677404	1.03
amino acid transporter system A1	NM_030674	1.03
hypothetical protein FLJ12820	NM_022451	1.03
serologically defined colon cancer antigen 1	NM_004713	1.03
RalGDS-like gene	AF186779	1.03
ciliary neurotrophic factor receptor	NM_001842	1.03
CGI-101 protein	NM_016041	1.03
hypothetical protein FLJ22242	NM_025070	1.03
megakaryoblastic leukemia (translocation) 1	AB037859	1.03
hypothetical protein FLJ11838	NM_024664	1.03
gb:M32577.1 /DEF=Human MHC HLA-DQ beta mRNA, complete cds. /FEA=mRNA /GEN=HLA-DQB1 /DB_XREF=gi:188194 /FL=gb:M32577.1	M32577	1.03
malonyl-CoA decarboxylase	NM_012213	1.03
rhodopsin (opsin 2, rod pigment) (retinitis pigmentosa 4, autosomal dominant)	AA058836	1.03
uncharacterized hypothalamus protein HCDASE	NM_018479	1.03
ATP-binding cassette, sub-family C (CFTR/MRP), member 4	AI948503	1.03
pancreatic polypeptide receptor 1	U42387	1.03
Consensus includes gb:AL359602.1 /DEF=Homo sapiens mRNA; cDNA DKFZp547D146 (from clone DKFZp547D146). /FEA=mRNA /DB_XREF=gi:8655670 /UG=Hs.306514 Homo sapiens mi	AL359602	1.03
Consensus includes gb:AW873544 /FEA=EST /DB_XREF=gi:8007597 /DB_XREF=est:ho62a01.x1 /CLONE=IMAGE:3041928 /UG=Hs.272500 Human mRNA for T cell receptor, clone IGRA17	AW873544	1.03
cytochrome P450, subfamily XXIA (steroid 21-hydroxylase, congenital adrenal hyperplasia), polypeptide 2	M17252	1.03
ATP synthase, H+ transporting, mitochondrial F1 complex, gamma polypeptide 1	AV711183	1.03
hypothetical protein FLJ10052	NM_017982	1.03
hypothetical protein FLJ14069	NM_025023	1.03
Consensus includes gb:AI339732 /FEA=EST /DB_XREF=gi:4076659 /DB_XREF=est:qq42f05.x1 /CLONE=IMAGE:1935201 /UG=Hs.309763 ESTs, Weakly similar to ALU7_HUMAN ALU SUBFA	AI339732	1.03
claudin 17	NM_012131	1.03
serologically defined breast cancer antigen NY-BR-20	AW025529	1.03
gb:NM_023939.1 /DEF=Homo sapiens hypothetical protein MGC2752 (MGC2752), mRNA. /FEA=mRNA /GEN=MGC2752 /PROD=hypothetical protein MGC2752 /DB_XREF=gi:13027609 /UG:	NM_023939	1.03
spinal cord-derived growth factor-B	NM_025208	1.03

CDC2-related protein kinase 7	NM_016507	1.03
aryl hydrocarbon receptor nuclear translocator-like	AB000815	1.03
hydroxyprostaglandin dehydrogenase 15-(NAD)	J05594	1.03
KIAA0710 gene product	NM_014871	1.03
hypothetical protein FLJ20254	NM_017727	1.03
hypothetical protein FLJ12552	BE856374	1.03
nuclear transcription factor, X-box binding 1	AF332009	1.03
p21(CDKN1A)-activated kinase 4	NM_005884	1.03
beta-carotene 15,15'-dioxygenase	NM_017429	1.03
DiGeorge syndrome critical region gene 6	NM_005675	1.03
SRY (sex determining region Y)-box 1	NM_005986	1.03
hypothetical protein FLJ14129	NM_030895	1.03
cyclin M1	NM_020348	1.03
putative G-protein coupled receptor	AK021758	1.03
melanoma-associated chondroitin sulfate proteoglycan-like	AF332228	1.03
KIAA0682 gene product	NM_014852	1.03
endothelin type b receptor-like protein 2	NM_004767	1.03
osmosis responsive factor	NM_012382	1.03
heterogeneous nuclear ribonucleoprotein A0	BE966599	1.03
leukocyte receptor cluster (LRC) member 4	BC006309	1.03
KIAA0239 protein	NM_015288	1.03
HBV pX associated protein-8	NM_016578	1.03
chromosome 1 open reading frame 28	NM_024529	1.03
KIAA0674 protein	AB014574	1.03
ubiquitin-conjugating enzyme E2A (RAD6 homolog)	NM_003336	1.03
Consensus includes gb:AI970157 /FEA=EST /DB_XREF=gi:5766983 /DB_XREF=est:wq89h08.x1 /CLONE=IMAGE:2479263 /UG=Hs.77631 glycine cleavage system protein H (aminomethyl c	AI970157	1.03
hypothetical protein Hy014	BF939727	1.03
advanced glycosylation end product-specific receptor	AB036432	1.03
KIAA0677 gene product	NM_014663	1.03
acrosomal vesicle protein 1	NM_001612	1.03
non-metastatic cells 3, protein expressed in	AL031718	1.03
SCAN domain-containing 1	AL109965	1.03
hypothetical protein FLJ12903	NM_022753	1.03
DKFZP586G011 protein	AK021613	1.03
phospholipase D1, phosphatidylcholine-specific	AI276230	1.03
growth hormone inducible transmembrane protein	AF131820	1.02
high-mobility group (nonhistone chromosomal) protein 1	NM_002128	1.02
leiomodin 1 (smooth muscle)	BC001755	1.02
tropomodulin	AI583393	1.02
adrenergic, alpha-1D-, receptor	M76446	1.02
Consensus includes gb:W48843 /FEA=EST /DB_XREF=gi 1337022 /DB_XREF=est:zc44a09.s1 /CLONE=IMAGE:325144 /UG=Hs.285814 sprouty (Drosophila) homolog 4 /FL=gb:AF227517.	W48843	1.02
melanoma antigen, family B, 1	NM_002363	1.02
histone acetyltransferase 1	NM_003642	1.02
Consensus includes gb:H95263 /FEA=EST /DB_XREF=gi 1102896 /DB_XREF=est:yu20e04.s1 /CLONE=IMAGE:234366 /UG=Hs.13431 Homo sapiens clone 24790 mRNA sequence	H95263	1.02
serum amyloid A4, constitutive	NM_006512	1.02
syntaxin 3A	AI002077	1.02
AHNAK nucleoprotein (desmoyokin)	BG287862	1.02
hypothetical protein FLJ12455	NM_022078	1.02
SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily a-like 1	NM_014140	1.02
KIAA0096 protein	AF226044	1.02
solute carrier family 25 (mitochondrial carrier; peroxisomal membrane protein, 34kD), member 17	AL049764	1.02
hypothetical protein FLJ23022	NM_025051	1.02
transcription factor-like 4	NM_013383	1.02
HIV-1 Rev binding protein	AI742626	1.02
KIAA0150 protein	D63484	1.02
breast cancer 1, early onset	NM_007295	1.02
protein phosphatase 1, regulatory (inhibitor) subunit 8	NM_002713	1.02
hypothetical protein FLJ23584	NM_024588	1.02
dynein, axonemal, light polypeptide 4	NM_005740	1.02
KIAA0426 gene product	NM_014724	1.02
TEA domain family member 1 (SV40 transcriptional enhancer factor)	AW771935	1.02
integrin, beta 1 (fibronectin receptor, beta polypeptide, antigen CD29 includes MDF2, MSK12)	AA149545	1.02
Consensus includes gb:AI659005 /FEA=EST /DB_XREF=gi:4762575 /DB_XREF=est:tu23e02.x1 /CLONE=IMAGE:2251898 /UG=Hs.4082 lectin, galactoside-binding, soluble, 8 (galectin 8) /	AI659005	1.02

nuclear factor related to kappa B binding protein	NM_006165	1.02
zinc finger protein 198	NM_003453	1.02
peptidase D	NM_000285	1.02
hypothetical protein MGC5297	NM_024091	1.02
Consensus includes gb:AW969913 /FEA=EST /DB_XREF=gi:8159757 /DB_XREF=est:EST381991 /UG=Hs.105121 ESTs	AW969913	1.02
hypothetical protein FLJ10233	NM_018034	1.02
dystrophia myotonica-protein kinase	M87313	1.02
nucleophosmin (nucleolar phosphoprotein B23, numatrin)	AA191576	1.02
splicing factor 3b, subunit 4, 49kD	BC004273	1.02
ribosomal protein L17	AL049597	1.02
Consensus includes gb:BF246115 /FEA=EST /DB_XREF=gi:11160133 /DB_XREF=est:601854068F1 /CLONE=IMAGE:4073921 /UG=Hs.8765 RNA helicase-related protein	BF246115	1.02
tripartite motif-containing 22	AA083478	1.02
hypothetical protein P15-2	AK023289	1.02
guanylate cyclase 1, soluble, beta 2	NM_004129	1.02
regulator of G-protein signalling 9	NM_003835	1.02
KIAA0475 gene product	NM_014864	1.02
KIAA1451 protein	AL049923	1.02
variable charge, X chromosome	AF167079	1.02
nuclear receptor subfamily 4, group A, member 1	NM_002135	1.02
KIAA0781 protein	AB018324	1.02
mitogen-activated protein kinase kinase kinase kinase 1	AA744529	1.02
DKFZP564G092 protein	NM_015601	1.02
mitochondrial ribosomal protein L20	NM_017971	1.02
melanoma antigen, family B, 3	NM_002365	1.02
serine (or cysteine) proteinase inhibitor, clade B (ovalbumin), member 2	NM_002575	1.02
chaperonin containing TCP1, subunit 6A (zeta 1)	BE737030	1.02
hypothetical protein MGC11335	NM_030819	1.02
v-jun sarcoma virus 17 oncogene homolog (avian)	BE327172	1.02
sulfotransferase family, cytosolic, 1A, phenol-preferring, member 2	NM_001054	1.02
septin 6	D50918	1.02
hypothetical protein FLJ13087	NM_025104	1.02
KIAA0864 protein	AK025604	1.02
hypothetical protein FLJ10656	NM_018170	1.02
SRY (sex determining region Y)-box 10	BC002824	1.02
uracil-DNA glycosylase	NM_003362	1.02
RYK receptor-like tyrosine kinase	X96588	1.02
insulin receptor-related receptor	J05046	1.02
signaling lymphocytic activation molecule	NM_003037	1.02
collagen, type VI, alpha 2	AL531750	1.02
Consensus includes gb:AI160126 /FEA=EST /DB_XREF=gi:3693506 /DB_XREF=est:qb51h08.x1 /CLONE=IMAGE:1703679 /UG=Hs.295901 KIAA0493 protein	AI160126	1.02
GA binding protein transcription factor, beta subunit 2 (47kD)	NM_002041	1.02
hypothetical protein FLJ22353	NM_024587	1.02
phosphatidylserine decarboxylase	AL096768	1.02
mitogen inducible 2	AW469573	1.02
hypothetical protein FLJ13213	NM_024755	1.02
GA binding protein transcription factor, alpha subunit (60kD)	D13318	1.02
homeo box D1	AI168371	1.02
stem cell growth factor; lymphocyte secreted C-type lectin	BC005810	1.02
PR domain containing 4	NM_012406	1.02
hypothetical protein FLJ12827	NM_024741	1.02
insulin promoter factor 1, homeodomain transcription factor	U30329	1.02
O-linked mannose beta1,2-N-acetylglucosaminyltransferase	NM_017739	1.02
PDZ domain protein (Drosophila inaD-like)	NM_005799	1.02
Homer, neuronal immediate early gene, 3	NM_004838	1.02
purine-rich element binding protein A	NM_005859	1.02
hypothetical protein FLJ20533	BC002748	1.02
DKFZP566B183 protein	BC002888	1.02
methionine adenosyltransferase II, alpha	BC001686	1.02
CGI-82 protein	NM_016026	1.02
cyclin E1	AI671049	1.02
hypothetical protein FLJ12661	NM_025138	1.02
dishevelled, dsh homolog 3 (Drosophila)	NM_004423	1.02
hypothetical protein FLJ12899	NM_024594	1.02

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ubiquitin specific protease 4 (proto-oncogene)	NM_003363	1.02
RAB3 GTPase-ACTIVATING PROTEIN	AI040009	1.02
Consensus includes gb:AF234255.1 /DEF=Homo sapiens clone KM36 immunoglobulin light chain variable region mRNA, partial cds. /FEA=mRNA /PROD=immunoglobulin light chain variable	AF234255	1.02
hypothetical protein FLJ23436	NM_024671	1.02
cytochrome c oxidase subunit VIIa polypeptide 2 (liver)	NM_001865	1.02
peptidylprolyl isomerase (cyclophilin)-like 2	U37220	1.02
Consensus includes gb:AW970944 /FEA=EST /DB_XREF=gi:8160789 /DB_XREF=est:EST383027 /UG=Hs.291839 ESTs	AW970944	1.02
solute carrier family 16 (monocarboxylic acid transporters), member 3	NM_004207	1.02
butyrophilin, subfamily 3, member A1	U90552	1.02
BTG family, member 3	AI765445	1.02
hypothetical protein FLJ12505	NM_024749	1.02
putative UDP-GalNAc:polypeptide N-acetylgalactosaminyltransferase T9	BE906572	1.02
sterile alpha and HEAT/Armadillo motif protein, ortholog of Drosophila	AI937297	1.02
pericentriolar material 1	AI924817	1.02
nuclear factor of kappa light polypeptide gene enhancer in B-cells inhibitor, beta	NM_002503	1.02
Consensus includes gb:AK000144.1 /DEF=Homo sapiens cDNA FLJ20137 fis, clone COL07137. /FEA=mRNA /DB_XREF=gi:7020040 /UG=Hs.274449 Homo sapiens cDNA FLJ20137 fis, clon	AK000144	1.02
KIAA1616 protein	AB046836	1.02
nuclear protein	AI357871	1.02
chromosome 11 open reading frame 13	H91083	1.02
bactericidal/permeability-increasing protein	NM_001725	1.02
KIAA1067 protein	BE646386	1.02
thymidine kinase 2, mitochondrial	NM_004614	1.02
fucosyltransferase 6 (alpha (1,3) fucosyltransferase)	H98825	1.02
estrogen receptor 2 (ER beta)	AF051428	1.02
BCL2-associated athanogene 5	AA457021	1.02
zinc finger protein 10 (KOX 1)	X52332	1.02
claudin 3	BE791251	1.02
islet cell autoantigen 1 (69kD)	W67992	1.02
kinesin family member 3B	NM_004798	1.02
glycogen synthase kinase 3 beta	BC000251	1.02
adaptor-related protein complex 3, delta 1 subunit	NM_003938	1.02
nuclear protein	AF273049	1.02
hypothetical protein FLJ23018	NM_024810	1.02
colony stimulating factor 2 (granulocyte-macrophage)	M11734	1.02
KIAA1196 protein	AI493119	1.02
Consensus includes gb:AI992251 /FEA=EST /DB_XREF=gi:5839156 /DB_XREF=est:ws41d12.x1 /CLONE=IMAGE:2499767 /UG=Hs.184581 ESTs	AI992251	1.02
G protein-coupled receptor kinase 5	NM_005308	1.02
hypothetical protein FLJ11328	NM_018391	1.02
CGI-146 protein	NM_016076	1.02
suppressor of Ty 6 homolog (S. cerevisiae)	NM_003170	1.02
susceptibility protein NSG-1	NM_016349	1.02
hypothetical protein FLJ21939 similar to 5-azacytidine induced gene 2	NM_022461	1.02
chromosome condensation 1-like	NM_001268	1.02
oxysterol binding protein-like 2	BC000296	1.02
ras-like protein	BE897886	1.02
clusterin (complement lysis inhibitor, SP-40,40, sulfated glycoprotein 2, testosterone-repressed prostate message 2, apolipoprotein J)	AI982754	1.02
retinol dehydrogenase 8 (all-trans)	NM_015725	1.02
Consensus includes gb:AC007204 /DEF=Homo sapiens chromosome 19, BAC 273239 (CIT-B-320G13) /FEA=CDS /DB_XREF=gi:4559317 /UG=Hs.284229 Homo sapiens chromosome 19, Ba	AC007204	1.02
single Ig IL-1R-related molecule	AA085764	1.02
KIAA0751 gene product	AF007156	1.02
solute carrier family 15 (H+/peptide transporter), member 2	BF223679	1.02
promyelocytic leukemia	NM_002675	1.02
latexin protein	NM_020169	1.02
vacuolar protein sorting 41 (yeast)	NM_014396	1.02
KIAA1028 protein	AI738802	1.02
leucine-rich repeat protein, neuronal 1	NM_002319	1.02
arylsulfatase D	NM_001669	1.02
apolipoprotein E	AI358867	1.02
Consensus includes gb:BE857715 /FEA=EST /DB_XREF=gi:10372018 /DB_XREF=est:7g46g03.x1 /CLONE=IMAGE:3309556 /UG=Hs.8982 ESTs, Highly similar to KIAA1395 protein H.sapien	BE857715	1.02
gb:NM_017421.1 /DEF=Homo sapiens methyltransferase COQ3 (COQ3), mRNA /FEA=mRNA /GEN=COQ3 /PROD=methyltransferase COQ3 /DB_XREF=gi:8393176 /FL=gb:NM_017421.1	NM_017421	1.02
clusterin (complement lysis inhibitor, SP-40,40, sulfated glycoprotein 2, testosterone-repressed prostate message 2, apolipoprotein J)	M25915	1.02
cullin 3	NM_003590	1.02
plasminogen	M74220	1.02

Consensus includes gb:AI798823 /FEA=EST /DB_XREF=gi:5364295 /DB_XREF=est:we93b05.x1 /CLONE=IMAGE:2348625 /UG=Hs.1050 pleckstrin homology, Sec7 and coiledcoil domains 1	AI798823	1.02
hypothetical protein FLJ20568	NM_017885	1.02
glucosaminyl (N-acetyl) transferase 1, core 2 (beta-1,6-N-acetylglucosaminyltransferase)	NM_001490	1.02
CGI-96 protein	AL589866	1.02
cleavage stimulation factor, 3' pre-RNA, subunit 1, 50kD	L02547	1.02
cold inducible RNA binding protein	NM_001280	1.02
GTF2I repeat domain-containing 1	NM_016328	1.02
ceroid-lipofuscinosis, neuronal 5	AI911687	1.02
gb:L14456.1 /DEF=Human Ig rearranged mu-chain gene V-N-D-N-J-region, complete cds. /FEA=mRNA /DB_XREF=gi:398497 /FL=gb:L14456.1	L14456	1.02
zinc finger protein 262	AA521508	1.02
KIAA0210 gene product	AI654161	1.02
hypothetical protein FLJ11101	NM_018322	1.02
KIAA0378 protein	Z38645	1.02
aristaless-like homeobox 4	NM_021926	1.02
KIAA0657 protein	AI633851	1.02
pancreatitis-associated protein	NM_002580	1.02
Consensus includes gb:AI097640 /FEA=EST /DB_XREF=gi:3445898 /DB_XREF=est:qb59a08.x1 /CLONE=IMAGE:1704374 /UG=Hs.66187 Homo sapiens clone 23700 mRNA sequence	AI097640	1.02
RAD51 homolog (RecA homolog, E. coli) (S. cerevisiae)	D14134	1.02
hypothetical protein LOC55565	NM_017530	1.02
transporter 1, ATP-binding cassette, sub-family B (MDR/TAP)	AI992187	1.02
BAI1-associated protein 3	AL031709	1.02
intracisternal A particle-promoted polypeptide	NM_005897	1.02
jagged 1 (Alagille syndrome)	U73936	1.02
5-hydroxytryptamine (serotonin) receptor 1D	NM_000864	1.02
Consensus includes gb:AK021690.1 /DEF=Homo sapiens cDNA FLJ11628 fis, clone HEMBA1004238. /FEA=mRNA /DB_XREF=gi:10432926 /UG=Hs.306610 Homo sapiens cDNA FLJ11628 fi	AK021690	1.02
KIAA0977 protein	BF002844	1.02
netrin 1	NM_004822	1.02
zinc finger protein 219	NM_016423	1.02
KIAA0599 protein	AI738980	1.02
hypothetical protein DKFZp434J037	NM_030952	1.02
EBNA1 binding protein 2	NM_006824	1.02
zinc finger protein 137 (clone pHZ-30)	NM_003438	1.02
frizzled homolog 8 (Drosophila)	AL121749	1.02
transcription termination factor, RNA polymerase I	AI632304	1.02
hypothetical protein FLJ14320	AK024382	1.02
tumor necrosis factor receptor superfamily, member 1A	NM_001065	1.01
KH-type splicing regulatory protein (FUSE binding protein 2)	NM_003685	1.01
CD3D antigen, delta polypeptide (TiT3 complex)	NM_000732	1.01
U6 snRNA-associated Sm-like protein	BC005938	1.01
proline-rich protein with nuclear targeting signal	AF279899	1.01
ribose 5-phosphate isomerase A (ribose 5-phosphate epimerase)	AI692341	1.01
Consensus includes gb:AK025097.1 /DEF=Homo sapiens cDNA: FLJ21444 fis, clone COL04437. /FEA=mRNA /DB_XREF=gi:10437544 /UG=Hs.306764 Homo sapiens cDNA: FLJ21444 fis, cl	AK025097	1.01
hypothetical protein FLJ10110	NM_017998	1.01
nuclear mitotic apparatus protein 1	NM_006185	1.01
CD84 antigen (leukocyte antigen)	NM_003874	1.01
Consensus includes gb:AL035297.1 /DEF=H.sapiens gene from PAC 747L4. /FEA=mRNA /PROD=hypothetical protein /DB_XREF=gi:4200248 /UG=Hs.119254 H.sapiens gene from PAC 747	AL035297	1.01
ems1 sequence (mammary tumor and squamous cell carcinoma-associated (p80/85 src substrate)	NM_005231	1.01
neuropeptide FF-amide peptide precursor	NM_003717	1.01
AU RNA binding protein/enoyl-Coenzyme A hydratase	NM_001698	1.01
nuclear domain 10 protein	BC004130	1.01
Consensus includes gb:AK024158.1 /DEF=Homo sapiens cDNA FLJ14096 fis, clone MAMMA1000752. /FEA=mRNA /DB_XREF=gi:10436469 /UG=Hs.303289 Homo sapiens cDNA FLJ14096 f	AK024158	1.01
KIAA0141 gene product	AI539425	1.01
fibulin 1	NM_001996	1.01
hypothetical protein FLJ13459	AF218012	1.01
XPMC2 protein	NM_020385	1.01
ATPase, Class V, type 10D	AI478147	1.01
placental protein 13	NM_013268	1.01
BUB1 budding uninhibited by benzimidazoles 1 homolog beta (yeast)	NM_001211	1.01
blepharophimosis, epicanthus inversus and ptosis, candidate 1	NM_021812	1.01
MGC6.1.1 protein	AB016900	1.01
CGI-29 protein	NM_015957	1.01
activin A receptor, type II	NM_001616	1.01
ribosomal protein L37	BF216701	1.01

K01391 B subtilis TrpE protein, TrpD protein, TrpC protein corresponding to nucleotides 1883-4400 of K01391 (-5, -M, -3 represent transcript regions 5 prime, Middle, and 3 prime respecti	K01391	1.01
melanocortin 3 receptor	NM_019888	1.01
lysophospholipase I	BG288007	1.01
phosphorylase kinase, beta	NM_000293	1.01
suppressor of potassium transport defect 3	AI655698	1.01
glucagon-like peptide 1 receptor	U01157	1.01
potassium voltage-gated channel, shaker-related subfamily, beta member 2	AF044253	1.01
actin, alpha 2, smooth muscle, aorta	NM_001613	1.01
POU domain, class 5, transcription factor 1	AF268613	1.01
transforming growth factor beta-stimulated protein TSC-22	AK027071	1.01
synaptophysin-like protein	NM_006754	1.01
G-protein coupled receptor SALPR; somatostatin and angiotensin-like peptide receptor	NM_016568	1.01
clathrin, light polypeptide (Lca)	NM_001833	1.01
ras-like protein	AW771590	1.01
hypothetical protein FLJ36421	AI133727	1.01
leukocyte immunoglobulin-like receptor, subfamily B (with TM and ITIM domains), member 6	NM_024318	1.01
ectonucleotide pyrophosphatase/phosphodiesterase 1	AU130282	1.01
hypothetical protein DKFZp564L2423	NM_030805	1.01
dual-specificity tyrosine-(Y)-phosphorylation regulated kinase 2	AI216690	1.01
RAB9, member RAS oncogene family	NM_004251	1.01
adrenal gland protein AD-004	NM_016283	1.01
dimethylarginine dimethylaminohydrolase 2	NM_013974	1.01
Consensus includes gb:AL031664 /DEF=Human DNA sequence from clone RP4-531H16 on chromosome 20p11.22-12. Contains the 3 part of the PCSK2 gene for proprotein convertase subti	AL031664	1.01
Consensus includes gb:D16471.1 /DEF=Human mRNA, Xq terminal portion. /FEA=mRNA /DB_XREF=gi:495099 /UG=Hs.121571 Human mRNA, Xq terminal portion	D16471	1.01
chloride channel 3	AA902971	1.01
Consensus includes gb:AL034396 /DEF=Human DNA sequence from clone 1158B12 on chromosome Xp11.21-11.4 Contains the ZXDA gene for X-linked duplicated Zinc finger A, and MYCL1	AL034396	1.01
nuclear receptor co-repressor 1	AB028970	1.01
general transcription factor IIE, polypeptide 1 (alpha subunit, 56kD)	NM_005513	1.01
nuclear transcription factor Y, alpha	AL031778	1.01
suppressor of S. cerevisiae gcr2	NM_007265	1.01
tumor necrosis factor receptor superfamily, member 5	NM_001250	1.01
LIM domain-containing preferred translocation partner in lipoma	AL044018	1.01
DNA segment, numerous copies, expressed probes (GS1 gene)	NM_012080	1.01
KIAA0470 gene product	AA126789	1.01
cholinergic receptor, nicotinic, alpha polypeptide 4	NM_000744	1.01
suppressor of K+ transport defect 1	AF195514	1.01
neuropathy target esterase	NM_006702	1.01
MAP-kinase activating death domain	AB002356	1.01
villin-like	BC004300	1.01
BCL2-antagonist/killer 1	NM_001188	1.01
glypican 1	AI354864	1.01
ubiquilin 3	NM_017481	1.01
palmitoyl-protein thioesterase 2	AF020543	1.01
phosphatidylinositol binding clathrin assembly protein	AL135735	1.01
MIRA interacting protein 5	NM_015700	1.01
gb:Z25422.1 /DEF=H.sapiens protein-serinethreonine kinase gene, complete CDS /FEA=mRNA /PROD=protein-serinethreonine kinase /DB_XREF=gi:405722 /FL=gb:Z25422.1	Z25422	1.01
retinol binding protein 4, plasma	NM_006744	1.01
calcium channel, voltage-dependent, alpha 1F subunit	NM_005183	1.01
pyrroline-5-carboxylate reductase 1	NM_006907	1.01
tubulin beta-5	BC002654	1.01
pregnancy-associated interferon	L25664	1.01
Consensus includes gb:AU146945 /FEA=EST /DB_XREF=gi:11008466 /DB_XREF=est:AU146945 /CLONE=HEMBB1001944 /UG=Hs.296687 Homo sapiens cDNA FLJ12040 fis, clone HEMBB	AU146945	1.01
hypothetical protein	BG167522	1.01
cadherin 16, KSP-cadherin	NM_004062	1.01
chromosome 2 open reading frame 2	NM_019063	1.01
Neuro-d4 (ral) homolog	NM_004647	1.01
Consensus includes gb:W88821 /FEA=EST /DB_XREF=gi:1404293 /DB_XREF=est:zh71d12.r1 /CLONE=IMAGE:417527 /UG=Hs.288924 Homo sapiens cDNA FLJ11392 fis, clone HEMBA100	W88821	1.01
cytochrome P450, subfamily IIB (phenobarbital-inducible), polypeptide 6	NM_000767	1.01
hypothetical protein FLJ10385	NM_018081	1.01
slit homolog 1 (Drosophila)	AB011537	1.01
chromosome 22 open reading frame 2	NM_015373	1.01
enigma (LIM domain protein)	NM_005451	1.01
H2B histone family, member A	BC001131	1.01

Consensus includes gb:AL135960 /DEF=Human DNA sequence from clone RP1-18D14 on chromosome 1 Contains genes for SIL (TAL1 (SCL) interrupting locus), TAL1 (T-cell acute lymphoc	AL135960	1.01
DKFZp566D133 protein	BG436400	1.01
SP140 nuclear body protein	NM_007237	1.01
SH2-B homolog	AL049924	1.01
nuclear factor of kappa light polypeptide gene enhancer in B-cells 1 (p105)	M55643	1.01
sperm autoantigenic protein 17	NM_017425	1.01
clustarin (complement lysis inhibitor, SP-40,40, sulfated glycoprotein 2, testosterone-repressed prostate message 2, apolipoprotein J)	M25915	1.01
Consensus includes gb:AI081543 /FEA=EST /DB_XREF=gi:3418335 /DB_XREF=est:on04g12.x1 /CLONE=IMAGE:1555750 /UG=Hs.300861 ESTs, Highly similar to T08701 hypothetical prot	AI081543	1.01
spermatogenesis associated 2	U28164	1.01
zinc finger protein 184 (Kruppel-like)	AI811577	1.01
transforming, acidic coiled-coil containing protein 3	NM_006342	1.01
olfactory receptor, family 7, subfamily E, member 38 pseudogene	AW874308	1.01
nuclear factor of kappa light polypeptide gene enhancer in B-cells inhibitor, alpha	AI078167	1.01
caspase 9, apoptosis-related cysteine protease	U60521	1.01
hypothetical protein FLJ13046 similar to exportin 4	NM_022459	1.01
myotubularin related protein 8	AW014788	1.01
synuclein, alpha (non A4 component of amyloid precursor)	L36674	1.01
PTH-responsive osteosarcoma B1 protein	U87408	1.01
phorbolin (similar to apolipoprotein B mRNA editing protein)	NM_004900	1.01
cholinergic receptor, nicotinic, epsilon polypeptide	AI859060	1.01
protein phosphatase 2 (formerly 2A), catalytic subunit, beta isoform	NM_004156	1.01
nucleophosmin (nucleolar phosphoprotein B23, numatrin)	BC002398	1.01
huntingtin interacting protein-1-related	AB013384	1.01
solute carrier family 28 (sodium-coupled nucleoside transporter), member 2	NM_004212	1.01
KIAA0671 gene product	AL136896	1.01
membrane-bound transcription factor protease, site 2	U73479	1.01
oxysterol binding protein 2	NM_030758	1.01
cardiac-specific homeo box	NM_004387	1.01
KIAA0453 protein	BF111962	1.01
Fanconi anemia, complementation group F	NM_022725	1.01
zinc finger protein 174	NM_003450	1.01
tripartite motif-containing 31	X81006	1.01
Consensus includes gb:R48779 /FEA=EST /DB_XREF=gi:810805 /DB_XREF=est:yj69c07.s1 /CLONE=IMAGE:153996 /UG=Hs.293845 ESTs, Weakly similar to B32688 beta-galactosidase-rel	R48779	1.01
sphingomyelin phosphodiesterase 1, acid lysosomal (acid sphingomyelinase)	M81780	1.01
early growth response 3	NM_004430	1.01
lymphocyte antigen 117	AF000424	1.01
solute carrier family 24 (sodium/potassium/calcium exchanger), member 3	NM_020689	1.01
neurofibromin 2 (bilateral acoustic neuroma)	BC003112	1.01
hypothetical protein FLJ10120	NM_018001	1.01
Consensus includes gb:W85912 /FEA=EST /DB_XREF=gi:1398341 /DB_XREF=est:zh52b12.s1 /CLONE=IMAGE:415679 /UG=Hs.323470 Homo sapiens cDNA: FLJ22106 fis, clone HEP17675	W85912	1.01
Consensus includes gb:AF005081.1 /DEF=Homo sapiens skin-specific protein (xp32) mRNA, partial cds. /FEA=mRNA /GEN=xp32 /PROD=skin-specific protein /DB_XREF=gi:2589189 /UG=	AF005081	1.01
hypothetical protein DKFZp586E1923	NM_020425	1.01
Consensus includes gb:AI246730 /FEA=EST /DB_XREF=gi:3842127 /DB_XREF=est:qk40b01.x1 /CLONE=IMAGE:1871401 /UG=Hs.126901 Homo sapiens mRNA full length insert cDNA clon	AI246730	1.01
Fzr1 protein	AA905473	1.01
staufen, RNA binding protein, homolog 2 (Drosophila)	NM_014393	1.01
KIAA0876 protein	AK026040	1.01
Misshapen/NIK-related kinase	AB041926	1.01
hypothetical protein FLJ20287	NM_017746	1.01
DKFZP434D193 protein	AK000323	1.01
hypothetical protein FLJ20342	NM_017774	1.01
solute carrier family 7 (cationic amino acid transporter, y+ system), member 6	BG230586	1.01
stannin	NM_003498	1.01
activating transcription factor 6	NM_007348	1.01
neogenin homolog 1 (chicken)	NM_002499	1.01
SP110 nuclear body protein	AA969194	1.01
KIAA1033 protein	AK001728	1.01
Consensus includes gb:AW081113 /FEA=EST /DB_XREF=gi:6036265 /DB_XREF=est:xc29c08.x1 /CLONE=IMAGE:2585678 /UG=Hs.18368 DKFZP564B0769 protein	AW081113	1.01
cytochrome c oxidase subunit VIIb	NM_001866	1.01
eukaryotic translation initiation factor 3, subunit 3 (gamma, 40kD)	NM_003756	1.01
islet-specific glucose-6-phosphatase catalytic subunit-related protein	NM_021176	1.01
plasminogen activator, urokinase receptor	AY029180	1.01
Fas (TNFRSF6)-associated via death domain	NM_003824	1.01
Mdm2, transformed 3T3 cell double minute 2, p53 binding protein (mouse)	NM_002392	1.01

transcription factor-like 1	NM_005997	1.01
Consensus includes gb:AF035278.1 /DEF=Homo sapiens clone 23676 mRNA sequence. /FEA=mRNA /DB_XREF=gi:2661027 /UG=Hs.100841 Homo sapiens clone 23676 mRNA sequence	AF035278	1.01
interleukin 17B	NM_014443	1.01
mitochondrial ribosomal protein L18	NM_014161	1.01
growth arrest-specific 1	AW611727	1.01
Consensus includes gb:AL390153.1 /DEF=Homo sapiens mRNA; cDNA DKFZp762G113 (from clone DKFZp762G113). /FEA=mRNA /DB_XREF=gi:9368841 /UG=Hs.208339 Homo sapiens mI	AL390153	1.01
nuclear factor of activated T-cells, cytoplasmic, calcineurin-dependent 1	U08015	1.01
KIAA1654 protein	AB051441	1.01
sentrin/SUMO-specific protease	NM_020654	1.01
actin related protein 2/3 complex, subunit 2 (34 kD)	NM_005731	1.01
cyclin T2	AV681875	1.01
hypothetical protein MGC2821	BC000769	1.01
KIAA1155 protein	AB032981	1.01
centaurin, delta 2	AB018325	1.01
bridging integrator 1	NM_004305	1.01
RecQ protein-like 5	AL136869	1.01
adaptor-related protein complex 1, sigma 2 subunit	NM_003916	1.01
amyloid beta precursor protein (cytoplasmic tail) binding protein 2	AV681579	1.00
zinc finger protein 278	AI807017	1.00
CGI-111 protein	NM_016048	1.00
Consensus includes gb:AI814252 /FEA=EST /DB_XREF=gi:5425467 /DB_XREF=est:wj70g04.x1 /CLONE=IMAGE:2408214 /UG=Hs.194051 Homo sapiens mRNA; cDNA DKFZp5668213 (fron	AI814252	1.00
transmembrane 4 superfamily member 7	BC000389	1.00
U3 snoRNP-associated 55-kDa protein	NM_004704	1.00
zinc/iron regulated transporter-like	NM_014437	1.00
immunoglobulin lambda joining 3	AB014341	1.00
GREB1 protein	AF245389	1.00
aldehyde dehydrogenase 7 family, member A1	BC002515	1.00
Consensus includes gb:AK024543.1 /DEF=Homo sapiens cDNA: FLJ20890 fis, clone ADKA03323. /FEA=mRNA /DB_XREF=gi:10436848 /UG=Hs.306686 Homo sapiens cDNA: FLJ20890 fis,	AK024543	1.00
Consensus includes gb:AA581879 /FEA=EST /DB_XREF=gi:2360557 /DB_XREF=est:nn43g06.s1 /CLONE=IMAGE:1086682 /UG=Hs.66191 Homo sapiens clone 24675 mRNA sequence	AA581879	1.00
Cluster Incl. AI057637:oy31h06.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-1667483 /clone_end=3 /gb=AI057637 /gi=3331503 /ug=Hs.21305 /len=599	AI057637	1.00
hypothetical protein FLJ22559	NM_024928	1.00
cadherin 4, type 1, R-cadherin (retinal)	NM_001794	1.00
phosphatidic acid phosphatase type 2B	AA628586	1.00
CCR4-NOT transcription complex, subunit 4	NM_013316	1.00
hypothetical protein MGC4365	BC003654	1.00
REV1-like (yeast)	NM_016316	1.00
KIAA0356 gene product	AI002220	1.00
hypothetical protein FLJ22555	NM_024520	1.00
ATP synthase, H+ transporting, mitochondrial F0 complex, subunit c (subunit 9), isoform 2	NM_005176	1.00
tyrosine 3-monooxygenase/tryptophan 5-monooxygenase activation protein, theta polypeptide	BF033313	1.00
cystatin B (cystatin-related epididymal specific)	NM_005492	1.00
homeo box A7	AF026397	1.00
uncharacterized bone marrow protein BM036	NM_018453	1.00
Consensus includes gb:AI807507 /FEA=EST /DB_XREF=gi:5394073 /DB_XREF=est:wf48e10.x1 /CLONE=IMAGE:2358858 /UG=Hs.126496 Homo sapiens chromosome 19, cosmid R31343	AI807507	1.00
hypothetical protein FLJ21324	AW449022	1.00
hypothetical protein MGC5528	NM_024094	1.00
KIAA0653 protein, B7-like protein	AL355690	1.00
Consensus includes gb:BF195104 /FEA=EST /DB_XREF=gi:11081627 /DB_XREF=est:7n15a07.x1 /CLONE=IMAGE:3564565 /UG=Hs.56729 lymphocyte-specific protein 1	BF195104	1.00
Consensus includes gb:AI346431 /FEA=EST /DB_XREF=gi:4083637 /DB_XREF=est:qp50g09.x1 /CLONE=IMAGE:1926496 /UG=Hs.75188 wee1+ (S. pombe) homolog	AI346431	1.00
trinucleotide repeat containing 3	AL541302	1.00
kinesin family member 3A	NM_007054	1.00
plasminogen activator, urokinase receptor	AC005525	1.00
Consensus includes gb:AL137474.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434E2423 (from clone DKFZp434E2423). /FEA=mRNA /DB_XREF=gi:6808085 /UG=Hs.244772 Homo sapiens	AL137474	1.00
NFS1 nitrogen fixation 1 (S. cerevisiae)	NM_021100	1.00
retinal outer segment membrane protein 1	NM_000327	1.00
Consensus includes gb:AL049037 /FEA=EST /DB_XREF=gi:4728346 /DB_XREF=est:DKFZp434P1518_r1 /CLONE=DKFZp434P1518 /UG=Hs.323833 syntaphilin	AL049037	1.00
KIAA0582 protein	NM_015147	1.00
like mouse brain protein E46	AF119662	1.00
solute carrier family 16 (monocarboxylic acid transporters), member 3	AA853175	1.00
chromosome 17 open reading frame 1A	U43383	1.00
chromosome 1 open reading frame 2	NM_006589	1.00
hypothetical protein MGC2508	NM_024327	1.00
hypothetical protein DKFZp762E1312	NM_018410	1.00

presenilin 1 (Alzheimer disease 3)	NM_007319	1.00
KIAA0871 protein	AI871641	1.00
actin-like 7B	NM_006686	1.00
folate transporter/carrier	NM_030780	1.00
olfactory receptor, family 2, subfamily N, member 1 pseudogene	AL022727	1.00
chorionic somatomammotropin hormone 2	NM_022646	1.00
inositol 1,4,5-triphosphate receptor, type 2	AA834576	1.00
TNF receptor-associated factor 5	NM_004619	1.00
HUG1 gene	NM_020527	1.00
suppressor of white apricot homolog 2	NM_007056	1.00
RAB30, member RAS oncogene family	NM_014488	1.00
ATP-binding cassette, sub-family F (GCN20), member 2	AF261091	1.00
synaptogyrin 1	BC000731	1.00
arylsulfatase A	U62317	1.00
tolloid-like 1	AI769310	1.00
Consensus includes gb:AI923972 /FEA=EST /DB_XREF=gi:5659936 /DB_XREF=est:wn49c10.x1 /CLONE=IMAGE:2448786 /UG=Hs.127723 Homo sapiens mRNA; cDNA DKFZp586J211 (from	AI923972	1.00
found in inflammatory zone 3	NM_020415	1.00
casein kinase 1, alpha 1	BF510713	1.00
Consensus includes gb:AI692428 /FEA=EST /DB_XREF=gi:4969768 /DB_XREF=est:wd69c12.x1 /CLONE=IMAGE:2336854 /UG=Hs.184019 Homo sapiens clone 23551 mRNA sequence	AI692428	1.00
hypothetical protein FLJ23591	NM_025076	1.00
UDP-Gal:betaGlcNAc beta 1,4- galactosyltransferase, polypeptide 6	NM_004775	1.00
N-acetyltransferase 1 (arylamine N-acetyltransferase)	NM_000662	1.00
Siah-interacting protein	AF057356	1.00
trinucleotide repeat containing 11 (THR-associated protein, 230 kDa subunit)	BC004354	1.00
putative DNA binding protein	AL117354	1.00
prostaglandin-endoperoxide synthase 1 (prostaglandin G/H synthase and cyclooxygenase)	S36219	1.00
PDZ-LIM protein mystique	NM_021630	1.00
paraoxonase 3	L48516	1.00
apoptosis inhibitor 5	AF229254	1.00
proteasome (prosome, macropain) subunit, alpha type, 4	NM_002789	1.00
actin-like 7A	NM_006687	1.00
formyltetrahydrofolate dehydrogenase	NM_012190	1.00
hypothetical protein DKFZp547P082	AL512728	1.00
sialyltransferase 4B (beta-galactosidase alpha-2,3-sialytransferase)	NM_006927	1.00
KIAA0349 protein	AB002347	1.00
sphingosine kinase type 2 isoform	AA485440	1.00
Consensus includes gb:AV691323 /FEA=EST /DB_XREF=gi:10293186 /DB_XREF=est:AV691323 /CLONE=GKCEWF11 /UG=Hs.2056 UDP glycosyltransferase 1 family, polypeptide A9	AV691323	1.00
chorionic somatomammotropin hormone-like 1	NM_022560	1.00
KIAA0265 protein	AF277177	1.00
glutamate receptor, ionotropic, kainate 5	S40369	1.00
ADP-ribosylation factor domain protein 1, 64kD	AF230398	1.00
macrophage receptor with collagenous structure	NM_006770	1.00
uridine phosphorylase	NM_003364	1.00
ketohexokinase (fructokinase)	BC006233	1.00
Consensus includes gb:AL389886 /DEF=Human DNA sequence from clone RP1-237C24 on chromosome 20 Contains the 3 end of the SLC23A1 gene encoding solute carrier family 23 membe	AL389886	1.00
three prime repair exonuclease 1	BC002903	1.00
RNA helicase-related protein	NM_007372	1.00
Consensus includes gb:Z95624 /DEF=Human DNA sequence from cosmid U237H1 contains Ras like GTPase and ESTs /FEA=CDS /DB_XREF=gi:2117165 /UG=Hs.248013 Human DNA seque	Z95624	1.00
DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 3	NM_024005	1.00
DKFZP434H132 protein	NM_015492	1.00
NY-REN-24 antigen	AI913329	1.00
T-cell leukemia translocation altered gene	NM_022171	1.00
cystathionase (cystathionine gamma-lyase)	NM_001902	1.00
hypothetical protein FLJ13055	NM_022737	1.00
synovial sarcoma, X breakpoint 2	BC002818	1.00
platelet-activating factor acetylhydrolase, isoform Ib, alpha subunit (45kD)	L13386	1.00
TAR (HIV) RNA binding protein 1	NM_005646	1.00
lysyl-tRNA synthetase	NM_005548	1.00
hypothetical protein FLJ12886	NM_019108	1.00
gb:AF130075.1 /DEF=Homo sapiens clone FLB9413 PRO2532 mRNA, complete cds. /FEA=mRNA /PROD=PRO2532 /DB_XREF=gi:11493455 /UG=Hs.306961 Homo sapiens clone FLB9413	AF130075	1.00
adenylate cyclase 3	AF033861	1.00
laminin, alpha 4	U77706	1.00
deafness, autosomal dominant 5	NM_004403	1.00

FIGURE 4D.

Gene	NCBI Accession Number	FOLD DECREASE
ribosomal protein S11	BF680255	22.06
ribosomal protein, large P2	BC005354	15.91
Consensus includes gb:AI249377.1 /DEF=Homo sapiens partial mRNA for human Ig lambda light chain variable region, clone MB91 (331 bp). /FEA=mRNA /GEN=IGLV /PROD=immunoglobu	AI249377	15.77
calreticulin	AI378706	14.14
ribosomal protein S19	BC000023	14.00
Consensus includes gb:D84143.1 /DEF=Human immunoglobulin (mAb59) light chain V region mRNA, partial sequence. /FEA=mRNA /PROD=immunoglobulin light chain V-J region /DB_XREF	D84143	13.82
Consensus includes gb:AA292281 /FEA=EST /DB_XREF=gi:1940261 /DB_XREF=est:zt51b03.s1 /CLONE=IMAGE:725837 /UG=Hs.181307 H3 histone, family 3A	AA292281	13.65
ribosomal protein L27a	BE737027	13.26
Consensus includes gb:AW302047 /FEA=EST /DB_XREF=gi:6711724 /DB_XREF=est:xr52f08.x1 /CLONE=IMAGE:2763783 /UG=Hs.76230 ribosomal protein S10	AW302047	13.16
ribosomal protein S20	AF113008	10.69
Consensus includes gb:L48784 /DEF=050 Homo sapiens cDNA /FEA=mRNA /DB_XREF=gi:1066715 /UG=Hs.182426 ribosomal protein S2	L48784	8.01
J04423 E coli bioB gene biotin synthetase (-5, -M, -3 represent transcript regions 5 prime, Middle, and 3 prime respectively)	J04423	6.76
calreticulin	AA910371	6.64
Escherichia coli /REF=J04423 /DEF=E coli bioB gene biotin synthetase corresponding to nucleotides 2393-2682 of J04423 /LEN=1114 (-5, -M, -3 represent transcript regions 5 prime, Mid	J04423	6.36
Cluster Incl. AI201594:qc02h12.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-1708487 /clone_end=3 /gb=AI201594 /gi=3754200 /ug=Hs.239333 /len=591	AI201594	6.20
ribosomal protein L27	BE312027	5.96
ribosomal protein L37a	BE857772	5.88
J04423 E coli bioB gene biotin synthetase (-5, -M, -3 represent transcript regions 5 prime, Middle, and 3 prime respectively)	J04423	5.80
Escherichia coli /REF=J04423 /DEF=E coli bioB gene biotin synthetase corresponding to nucleotides 2071-2304 of J04423 /LEN=1114 (-5, -M, -3 represent transcript regions 5 prime, Mid	J04423	5.77
Escherichia coli /REF=J04423 /DEF=E coli bioB gene biotin synthetase corresponding to nucleotides 2772-3004 of J04423 /LEN=1114 (-5, -M, -3 represent transcript regions 5 prime, Mid	J04423	5.72
J04423 E coli bioB gene biotin synthetase (-5, -M, -3 represent transcript regions 5 prime, Middle, and 3 prime respectively)	J04423	5.57
immunoglobulin lambda locus	AF043586	5.53
J04423 E coli bioC protein (-5 and -3 represent transcript regions 5 prime and 3 prime respectively)	J04423	5.35
ribosomal protein L38	AW303136	4.99
Consensus includes gb:AI239383.1 /DEF=Homo sapiens mRNA for immunoglobulin heavy chain variable region, ID 31. /FEA=mRNA /GEN=IGHV /PROD=immunoglobulin heavy chain variable	AI239383	4.96
Consensus includes gb:AI345238 /FEA=EST /DB_XREF=gi:4082444 /DB_XREF=est:tb81b07.x1 /CLONE=IMAGE:2060725 /UG=Hs.111334 ferritin, light polypeptide	AI345238	4.95
Escherichia coli /REF=J04423 /DEF=E coli bioC protein corresponding to nucleotides 4257-4573 of J04423 /LEN=777 (-5 and -3 represent transcript regions 5 prime and 3 prime respectiv	J04423	4.76
hypothetical protein FLJ21034	NM_024940	4.69
Escherichia coli /REF=J04423 /DEF=E coli bioC protein corresponding to nucleotides 4609-4883 of J04423 /LEN=777 (-5 and -3 represent transcript regions 5 prime and 3 prime respectiv	J04423	4.64
ribosomal protein L38	BC000603	4.47
killer cell immunoglobulin-like receptor, three domains, long cytoplasmic tail, 2	X93596	4.39
PTPRF interacting protein, binding protein 1 (liprin beta 1)	N35896	4.31
hypothetical protein FLJ12619	BE465032	4.30
C-terminal binding protein 1	BF984434	4.29
Consensus includes gb:BF942161 /FEA=EST /DB_XREF=gi:12359481 /DB_XREF=est:nae87g10.x1 /CLONE=IMAGE:4118994 /UG=Hs.302797 ESTs	BF942161	4.27
Consensus includes gb:AF044592 /DEF=Homo sapiens lymphocyte-predominant Hodgkins disease case 4 immunoglobulin heavy chain gene, variable region, partial cds /FEA=CDS /DB_XREI	AF044592	4.24
hypothetical protein FLJ12985	NM_024924	4.24
thiopurine S-methyltransferase	U12387	4.22
J04423 E coli bioD gene dethiobiotin synthetase (-5 and -3 represent transcript regions 5 prime and 3 prime respectively)	J04423	4.19
J04423 E coli bioC protein (-5 and -3 represent transcript regions 5 prime and 3 prime respectively)	J04423	4.17
Consensus includes gb:AL050122.1 /DEF=Homo sapiens mRNA; cDNA DKFZp586E121 (from clone DKFZp586E121). /FEA=mRNA /DB_XREF=gi:4884330 /UG=Hs.274511 Homo sapiens mF	AL050122	4.11
Consensus includes gb:AW301806 /FEA=EST /DB_XREF=gi:6711483 /DB_XREF=est:xr56e11.x1 /CLONE=IMAGE:2764172 /UG=Hs.150551 ESTs, Weakly similar to ALU1_HUMAN ALU SUBFI	AW301806	4.10
thromboxane A2 receptor	NM_001060	4.08
ADP-ribosylation factor 6	AA243143	4.02
PCTAIRE protein kinase 1	NM_006201	3.95
ITV1 gene	AF116615	3.90
hypothetical protein PRO1843	NM_018507	3.86
dystrophia myotonica-containing WD repeat motif	L19267	3.79
emopamil binding protein (sterol isomerase)	N58493	3.78
Escherichia coli /REF=J04423 /DEF=E coli bioD gene dethiobiotin synthetase corresponding to nucleotides 5312-5559 of J04423 /LEN=676 (-5 and -3 represent transcript regions 5 prime	J04423	3.77
Escherichia coli /REF=J04423 /DEF=E coli bioD gene dethiobiotin synthetase corresponding to nucleotides 5024-5244 of J04423 /LEN=676 (-5 and -3 represent transcript regions 5 prime	J04423	3.74
gb:AF090895.1 /DEF=Homo sapiens clone HQ0117 PRO0117 mRNA, complete cds. /FEA=mRNA /PROD=PRO0117 /DB_XREF=gi:6690166 /UG=Hs.283919 Homo sapiens clone HQ0117 PR	AF090895	3.71
GM2 ganglioside activator protein	X61094	3.70
putative protein O-mannosyltransferase	NM_013382	3.69
Consensus includes gb:BF973387 /FEA=EST /DB_XREF=gi:12340602 /DB_XREF=est:602242353F1 /CLONE=IMAGE:4330861 /UG=Hs.305989 Human DNA sequence from clone RP3-483K	BF973387	3.69
M10098 Human 18S rRNA sequence, length 1969 bases, middle target bases 647-1292	M10098	3.69
aminopeptidase puromycin sensitive	BG153399	3.65
gb:Z25437.1 /DEF=H.sapiens protein-tyrosine kinase gene, complete CDS /FEA=mRNA /PROD=protein-tyrosine kinase /DB_XREF=gi:405752 /FL=gb:Z25437.1	Z25437	3.63
Consensus includes gb:AK026825.1 /DEF=Homo sapiens cDNA: FLJ23172 fis, clone LNG10005. /FEA=mRNA /DB_XREF=gi:10439771 /UG=Hs.306885 Homo sapiens cDNA: FLJ23172 fis, cl	AK026825	3.63
ubiquitously transcribed tetratricopeptide repeat gene, Y chromosome	NM_007125	3.61

Consensus includes gb:AI732770 /FEA=EST /DB_XREF=gi:5053883 /DB_XREF=est:zx78d05.x5 /CLONE=IMAGE:809865 /UG=Hs.328688 ESTs, Moderately similar to ALU7_HUMAN ALU SUE	AI732770	3.57
acidic epididymal glycoprotein-like 1	X95238	3.55
hypothetical protein FLJ10357	NM_018071	3.51
J04423 E coli bioD gene dethiobiotin synthetase (-5 and -3 represent transcript regions 5 prime and 3 prime respectively)	J04423	3.48
Consensus includes gb:AV719355 /FEA=EST /DB_XREF=gi:10816507 /DB_XREF=est:AV719355 /CLONE=GLCEMB06 /UG=Hs.97109 ESTs	AV719355	3.44
KIAA1655 protein	AB051442	3.41
Consensus includes gb:AA890010 /FEA=EST /DB_XREF=gi:3016889 /DB_XREF=est:aj89h08.s1 /CLONE=IMAGE:1403679 /UG=Hs.50785 SEC22, vesicle trafficking protein (S. cerevisiae)-lil	AA890010	3.41
hypothetical protein FLJ21603	NM_024762	3.41
protein kinase, interferon-inducible double stranded RNA dependent	NM_002759	3.40
hypothetical protein FLJ10298	NM_018050	3.40
translation initiation factor IF2	AB018284	3.39
collagen, type I, alpha 1	AJ743621	3.39
alanine-glyoxylate aminotransferase 2-like 1	NM_031279	3.37
PRO0478 protein	NM_014129	3.36
Consensus includes gb:AW971134 /FEA=EST /DB_XREF=gi:8160979 /DB_XREF=est:EST383221 /UG=Hs.292245 ESTs, Weakly similar to ALU1_HUMAN ALU SUBFAMILY J SEQUENCE CONTAI	AW971134	3.32
sialophorin (gpL115, leukosialin, CD43)	NM_003123	3.31
islet cell autoantigen 1 (69kD)	BC005922	3.30
ATPase, H+ transporting, lysosomal (vacuolar proton pump) non-catalytic accessory protein 1A (110/116kD)	NM_005177	3.30
protein phosphatase 2 (formerly 2A), regulatory subunit B" (PR 72), alpha isoform and (PR 130), beta isoform	AI760130	3.29
HIV-1 rev binding protein 2	AJ912583	3.28
sorting nexin 4	AA524345	3.26
aldehyde dehydrogenase 1 family, member B1	BC001619	3.26
Consensus includes gb:AW971415 /FEA=EST /DB_XREF=gi:8161260 /DB_XREF=est:EST383504 /UG=Hs.165337 ESTs	AW971415	3.25
Consensus includes gb:AK026484.1 /DEF=Homo sapiens cDNA: FLJ22831 fis, clone KAIA4161. /FEA=mRNA /DB_XREF=gi:10439356 /UG=Hs.321666 Homo sapiens cDNA: FLJ22831 fis, cl	AK026484	3.25
6 protein-coupled receptor 37 (endothelin receptor type B-like)	T16257	3.24
hypothetical protein FLJ14107	NM_025026	3.22
endomucin-1	NM_016241	3.22
pregnancy specific beta-1-glycoprotein 11	NM_002785	3.22
tyrosine 3-monooxygenase/tryptophan 5-monooxygenase activation protein, epsilon polypeptide	AA502643	3.21
Consensus includes gb:AI701156 /FEA=EST /DB_XREF=gi:4989056 /DB_XREF=est:we10f09.x1 /CLONE=IMAGE:2340713 /UG=Hs.6580 Homo sapiens cDNA: FLJ23227 fis, clone CAE00645	AI701156	3.20
Consensus includes gb:AK022473.1 /DEF=Homo sapiens cDNA FLJ12411 fis, clone MAMMA1002964. /FEA=mRNA /DB_XREF=gi:10433882 /UG=Hs.296722 Homo sapiens cDNA FLJ12411	AK022473	3.18
CDC5 cell division cycle 5-like (S. pombe)	AB007892	3.17
Consensus includes gb:BF573849 /FEA=EST /DB_XREF=gi:11647561 /DB_XREF=est:602132053F1 /CLONE=IMAGE:4271340 /UG=Hs.96343 ESTs, Weakly similar to ALUC_HUMAN !!!! ALU	BF573849	3.17
harakiri, BCL2 interacting protein (contains only BH3 domain)	U76376	3.17
cytochrome P450 isoform 4F12	NM_023944	3.17
gb:BC006361.1 /DEF=Homo sapiens, clone MGC:13137, mRNA, complete cds. /FEA=mRNA /PROD=Unknown (protein for MGC:13137) /DB_XREF=gi:13623508 /FL=gb:BC006361.1	BC006361	3.16
hypothetical protein FLJ22965	NM_022101	3.15
Consensus includes gb:AF222691.1 /DEF=Homo sapiens Alu repeat (LNX1) mRNA sequence. /FEA=mRNA /DB_XREF=gi:12655850 /UG=Hs.307008 Homo sapiens Alu repeat (LNX1) mRNA	AF222691	3.15
clone FLB3816	NM_016415	3.15
Consensus includes gb:AI524687 /FEA=EST /DB_XREF=gi:4438822 /DB_XREF=est:th12a07.x1 /CLONE=IMAGE:2118036 /UG=Hs.57969 phenylalanine-tRNA synthetase	AI524687	3.14
hypothetical protein FLJ20897	AI335509	3.12
PI-3-kinase-related kinase SMG-1	BE000837	3.11
Consensus includes gb:AI683552 /FEA=EST /DB_XREF=gi:4893734 /DB_XREF=est:tx67h02.x1 /CLONE=IMAGE:2274675 /UG=Hs.201605 ESTs, Moderately similar to ALU8_HUMAN ALU SU	AI683552	3.08
KIAA0729 protein	AK023845	3.07
KIAA1827 protein	AW474158	3.07
C-type (calcium dependent, carbohydrate-recognition domain) lectin, superfamily member 6	AF200738	3.04
SCAN domain-containing 2	AF244812	3.03
thyroid hormone receptor-associated protein, 95-kD subunit	BG339606	3.02
Consensus includes gb:AK024108.1 /DEF=Homo sapiens cDNA FLJ14046 fis, clone HEMBA1006461. /FEA=mRNA /DB_XREF=gi:10436406 /UG=Hs.142677 Homo sapiens cDNA FLJ14046 f	AK024108	3.02
tumor necrosis factor alpha-inducible cellular protein containing leucine zipper domains; Huntingtin interacting protein L; transcrption factor IIIA-interacting protein	NM_021980	3.02
Consensus includes gb:BG290532 /FEA=EST /DB_XREF=gi:13047560 /DB_XREF=est:602388395F1 /CLONE=IMAGE:4517206 /UG=Hs.11210 ESTs, Moderately similar to Z137_HUMAN ZIN	BG290532	3.01
mitochondrial ribosomal protein S12	R68573	3.00
P3ECSL	NM_022164	2.99
lymphoid blast crisis oncogene	AF127481	2.99
KIAA0653 protein, B7-like protein	AF289028	2.99
Consensus includes gb:AF035317.1 /DEF=Homo sapiens clone 23892 mRNA sequence. /FEA=mRNA /DB_XREF=gi:2661080 /UG=Hs.91916 Homo sapiens clone 23892 mRNA sequence	AF035317	2.99
early lymphoid activation protein	L22650	2.98
lipopolysaccharide specific response-68 protein	NM_018678	2.97
tissue factor pathway inhibitor (lipoprotein-associated coagulation inhibitor)	J03225	2.96
Consensus includes gb:AI734156 /FEA=EST /DB_XREF=gi:5055269 /DB_XREF=est:nc79d04.y5 /CLONE=IMAGE:783559 /UG=Hs.172895 ESTs	AI734156	2.96
Consensus includes gb:AW301937 /FEA=EST /DB_XREF=gi:6711614 /DB_XREF=est:xr85b03.x1 /CLONE=IMAGE:2766893 /UG=Hs.138036 ESTs	AW301937	2.95
immunoglobulin heavy constant mu	S55735	2.93
Consensus includes gb:X78262.1 /DEF=H.sapiens mRNA for TRE5. /FEA=mRNA /DB_XREF=gi:587440 /UG=Hs.302178 H.sapiens mRNA for TRE5	X78262	2.92

Consensus includes gb:R06655 /FEA=EST /DB_XREF=gi:757275 /DB_XREF=est:yf10e02.r1 /CLONE=IMAGE:126458 /UG=Hs.188518 ESTs, Moderately similar to AF078844 1 hqp0376 prot	R06655	2.91
somatostatin receptor 4	NM_001052	2.91
Consensus includes gb:AL163202 /DEF=Homo sapiens chromosome 21 segment HS21C002 /FEA=CDS /DB_XREF=gi:7717242 /UG=Hs.289121 Homo sapiens chromosome 21 segment HS2	AL163202	2.91
guanine nucleotide exchange factor for Rap1; M-Ras-regulated GEF	AI263837	2.90
hypothetical protein PRO0082	NM_018590	2.89
cathepsin S	BC002642	2.88
transducin (beta)-like 1	AA724134	2.88
Consensus includes gb:AK022303.1 /DEF=Homo sapiens cDNA FLJ12241 fis, clone MAMMA1001274. /FEA=mRNA /DB_XREF=gi:10433670 /UG=Hs.287503 Homo sapiens cDNA FLJ12241 f	AK022303	2.87
MAD, mothers against decapentaplegic homolog (Drosophila) interacting protein, receptor activation anchor	NM_007323	2.87
Consensus includes gb:AL050065.1 /DEF=Homo sapiens mRNA; cDNA DKFZp566M043 (from clone DKFZp566M043). /FEA=mRNA /DB_XREF=gi:4884295 /UG=Hs.212587 Homo sapiens m	AL050065	2.86
E74-like factor 4 (ets domain transcription factor)	NM_001421	2.85
hypothetical protein 384D8_6	BC000473	2.84
phosphodiesterase 10A	AF127480	2.84
PABP-interacting protein 2	AL043487	2.84
PRO1880 protein	NM_014104	2.83
serine protease inhibitor-like, with Kunitz and WAP domains 1 (eppin)	NM_020398	2.82
zinc finger protein 43 (HTF6)	AK022905	2.82
solute carrier family 4, sodium bicarbonate cotransporter, member 4	AF011390	2.82
rab3 GTPase-activating protein, non-catalytic subunit (150kD)	AK021928	2.81
carboxypeptidase N, polypeptide 2, 83kD	I05158	2.79
hypothetical protein FLJ12151	AK022213	2.79
neuronal thread protein	NM_014486	2.78
Consensus includes gb:AK021505.1 /DEF=Homo sapiens cDNA FLJ11443 fis, clone HEMBA1001330. /FEA=mRNA /DB_XREF=gi:10432701 /UG=Hs.297945 Homo sapiens cDNA FLJ11443 fi	AK021505	2.78
KIAA0889 protein	NM_015377	2.78
coactivator-associated arginine methyltransferase-1	AL529396	2.77
ribonuclease P, 40kD subunit	NM_006638	2.76
solute carrier family 21 (organic anion transporter), member 6	AB026257	2.75
putative N6-DNA-methyltransferase	NM_013240	2.75
Consensus includes gb:AL080160.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434M054 (from clone DKFZp434M054). /FEA=mRNA /DB_XREF=gi:5262622 /UG=Hs.274517 Homo sapiens m	AL080160	2.74
dihydrolipoamide branched chain transacylase (E2 component of branched chain keto acid dehydrogenase complex; maple syrup urine disease)	NM_001918	2.73
transforming, acidic coiled-coil containing protein 2	AF220152	2.73
prostate derived STE20-like kinase PSK	NM_016151	2.73
Consensus includes gb:AL157484.1 /DEF=Homo sapiens mRNA; cDNA DKFZp762M127 (from clone DKFZp762M127). /FEA=mRNA /DB_XREF=gi:7018527 /UG=Hs.22483 Homo sapiens mR	AL157484	2.73
dynamin 2	AK023207	2.73
sema domain, immunoglobulin domain (Ig), short basic domain, secreted, (semaphorin) 3A	NM_006080	2.72
G protein-coupled receptor kinase 6	BG423052	2.72
S100 calcium binding protein A11 (calgizzarin)	NM_005620	2.72
cytochrome c oxidase subunit Vb	AI557312	2.72
Epstein-Barr virus induced gene 3	NM_005755	2.71
Consensus includes gb:AW974816 /FEA=EST /DB_XREF=gi:8166019 /DB_XREF=est:EST386921 /UG=Hs.283517 ESTs, Weakly similar to ALU1_HUMAN ALU SUBFAMILY J SEQUENCE CONTAI	AW974816	2.71
bone morphogenetic protein receptor, type IB	D89675	2.71
prefoldin 4	NM_002623	2.70
hypothetical protein FLJ21032	NM_024906	2.70
glioma pathogenesis-related protein	U16307	2.70
epithelial membrane protein 3	NM_001425	2.69
Consensus includes gb:AK024958.1 /DEF=Homo sapiens cDNA: FLJ21305 fis, clone COL02124. /FEA=mRNA /DB_XREF=gi:10437382 /UG=Hs.287658 Homo sapiens cDNA: FLJ21305 fis, cl	AK024958	2.69
KRAB zinc finger protein KR1B	AK024789	2.69
putatative 28 kDa protein	AF349314	2.68
apolipoprotein L, 2	BC004395	2.68
Consensus includes gb:AL110201.1 /DEF=Homo sapiens mRNA; cDNA DKFZp586F1622 (from clone DKFZp586F1622). /FEA=mRNA /DB_XREF=gi:5817120 /UG=Hs.278435 Homo sapiens	AL110201	2.67
Consensus includes gb:R33964 /FEA=EST /DB_XREF=gi:789822 /DB_XREF=est:yh74c03.r1 /CLONE=IMAGE:135460 /UG=Hs.288681 Homo sapiens cDNA FLJ11022 fis, clone PLACE1003737	R33964	2.67
Consensus includes gb:AL137378.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434K1126 (from clone DKFZp434K1126). /FEA=mRNA /DB_XREF=gi:6807908 /UG=Hs.306455 Homo sapiens	AL137378	2.67
translation initiation factor IF2	BG261322	2.67
KIAA1045 protein	AB028968	2.66
hypothetical protein	NM_019069	2.66
hypothetical protein FLJ23185	NM_025056	2.66
gamma-aminobutyric acid (GABA) A receptor, alpha 5	BF966183	2.66
colony stimulating factor 2 receptor, alpha, low-affinity (granulocyte-macrophage)	L29349	2.66
N-myristoyltransferase 2	NM_004808	2.64
IM27 protein	NM_007003	2.64
Consensus includes gb:AK023911.1 /DEF=Homo sapiens cDNA FLJ13849 fis, clone THYRO1000865. /FEA=mRNA /DB_XREF=gi:10435992 /UG=Hs.181810 Homo sapiens cDNA FLJ13849 fi	AK023911	2.63
immunoglobulin heavy constant mu	S74639	2.63
gb:NM_024305.1 /DEF=Homo sapiens hypothetical protein MGC4278 (MGC4278), mRNA. /FEA=mRNA /GEN=MGC4278 /PROD=hypothetical protein MGC4278 /DB_XREF=gi:13236535 /UG=	NM_024305	2.63

hypothetical protein MGC2474	NM_023931	2.62
hypothetical protein FLJ21477	NM_025153	2.62
polymerase (DNA directed), eta	NM_006502	2.62
Consensus includes gb:AI126492 /FEA=EST /DB_XREF=gi:3595006 /DB_XREF=est:qd82h06.x1 /CLONE=IMAGE:1736027 /UG=Hs.104258 Homo sapiens mRNA, exon 1, 2, 3, 4, clone:RES4	AI126492	2.62
Consensus includes gb:BC005365.1 /DEF=Homo sapiens, clone IMAGE:3829438, mRNA, partial cds. /FEA=mRNA /PROD=Unknown (protein for IMAGE:3829438) /DB_XREF=gi:13529199 /L	BC005365	2.61
polymerase (RNA) II (DNA directed) polypeptide B (140kD)	AW770896	2.61
cAMP responsive element binding protein-like 1	U52696	2.61
neuropeptide Y receptor Y6 (pseudogene)	U59431	2.60
hexokinase 2	AI761561	2.60
Cluster Incl. AI949010:wq36a07.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-2473332 /clone_end=3 /gb=AI949010 /gi=5741320 /ug=Hs.104036 /len=457	AI949010	2.59
mannan-binding lectin serine protease 1 (C4/C2 activating component of Ra-reactive factor)	BC000587	2.59
SH3-domain GRB2-like 3	AF036269	2.59
thyroid hormone receptor interactor 11	BC002656	2.59
hypothetical protein PRO2849	NM_022335	2.58
decay accelerating factor for complement (CD55, Cromer blood group system)	BC001288	2.58
mitogen-activated protein kinase kinase 5	U71088	2.58
cofactor required for Sp1 transcriptional activation, subunit 2 (150kD)	AK023368	2.57
phosphatidylinositol 4-kinase, catalytic, beta polypeptide	U81802	2.56
HCF-binding transcription factor Zhangfei	AI206560	2.56
kangai 1 (suppression of tumorigenicity 6, prostate; CD82 antigen (R2 leukocyte antigen, antigen detected by monoclonal and antibody IA4))	NM_002231	2.56
DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 17 (72kD)	NM_030881	2.55
RAP2B, member of RAS oncogene family	NM_002886	2.55
eukaryotic translation initiation factor 4 gamma, 1	BE966878	2.54
KIAA0472 protein	AB007941	2.54
isocitrate dehydrogenase 1 (NADP+), soluble	NM_005896	2.54
CED-6 protein	AK023668	2.53
ATP-binding cassette, sub-family D (ALD), member 3	NM_002858	2.53
eukaryotic translation initiation factor 5A	AA393940	2.53
glycophorin E	NM_002102	2.53
artemis protein	NM_022487	2.52
inhibin, beta C	NM_005538	2.52
reversion-inducing-cysteine-rich protein with kazal motifs	AK022897	2.52
Consensus includes gb:AU147017 /FEA=EST /DB_XREF=gi:11008538 /DB_XREF=est:AU147017 /CLONE=HEMBB1002152 /UG=Hs.301905 Homo sapiens cDNA FLJ14080 fis, clone HEMBB	AU147017	2.52
Mediterranean fever	NM_000243	2.52
hypothetical protein FLJ20837	NM_017964	2.52
Consensus includes gb:AL049260.1 /DEF=Homo sapiens mRNA; cDNA DKFZp564E233 (from clone DKFZp564E233). /FEA=mRNA /DB_XREF=gi:4500007 /UG=Hs.302050 Homo sapiens mI	AL049260	2.51
Consensus includes gb:AK025724.1 /DEF=Homo sapiens cDNA: FLJ22071 fis, clone HEP11691. /FEA=mRNA /DB_XREF=gi:10438333 /UG=Hs.326248 Homo sapiens cDNA: FLJ22071 fis, cl	AK025724	2.51
coagulation factor V (proaccelerin, labile factor)	NM_000130	2.49
ring finger protein 2	NM_007212	2.49
ATPase, Class I, type 8B, member 1	BG252666	2.49
homeo box A10	AI375919	2.49
Consensus includes gb:AF043583.1 /DEF=Homo sapiens clone ASMneg1-b3 immunoglobulin lambda chain VI region, (IGL) mRNA, partial cds. /FEA=mRNA /GEN=IGL /PROD=immunoglobul	AF043583	2.49
nucleolar protein 4	NM_003787	2.48
macrophage stimulating, pseudogene 9	U28055	2.48
death-associated protein kinase 2	AK026801	2.48
ubiquitin specific protease 15	AF106069	2.48
Consensus includes gb:AW301806 /FEA=EST /DB_XREF=gi:6711483 /DB_XREF=est:xr56e11.x1 /CLONE=IMAGE:2764172 /UG=Hs.150551 ESTs, Weakly similar to ALU1_HUMAN ALU SUBF/	AW301806	2.47
Fc fragment of IgG, low affinity IIIb, receptor for (CD16)	I04162	2.47
Consensus includes gb:U43604.1 /DEF=Human unidentified mRNA, partial sequence. /FEA=mRNA /DB_XREF=gi:1171236 /UG=Hs.159901 Human unidentified mRNA, partial sequence	U43604	2.47
prostate differentiation factor	AF003934	2.47
X03453 Bacteriophage P1 cre recombinase protein (-5 and -3 represent transcript regions 5 prime and 3 prime respectively)	X03453	2.47
protein kinase, cAMP-dependent, regulatory, type II, alpha	BC002763	2.47
hypothetical protein FLJ10534	AK026565	2.46
Consensus includes gb:AW574933 /FEA=EST /DB_XREF=gi:7246472 /DB_XREF=est:UI-HF-BL0-abq-b-09-0-UI.s1 /CLONE=IMAGE:3057353 /UG=Hs.248844 ESTs, Weakly similar to ALU1_H	AW574933	2.46
nerve growth factor receptor (TNFR superfamily, member 16)	NM_002507	2.46
neuropilin 2	AA295257	2.46
erythrocyte membrane protein band 7.2 (stomatin)	AI537887	2.46
ladinin 1	U58994	2.46
Consensus includes gb:AK026493.1 /DEF=Homo sapiens cDNA: FLJ22840 fis, clone KAIA4709. /FEA=mRNA /DB_XREF=gi:10439366 /UG=Hs.287293 Homo sapiens cDNA: FLJ22840 fis, cl	AK026493	2.45
heterogeneous nuclear ribonucleoprotein H1 (H)	AV753392	2.45
adducin 2 (beta)	NM_017482	2.45
zinc finger protein 76 (expressed in testis)	NM_003427	2.45
Consensus includes gb:BE672313 /FEA=EST /DB_XREF=gi:10032854 /DB_XREF=est:7a59b10.x1 /CLONE=IMAGE:3223003 /UG=Hs.34054 Homo sapiens cDNA: FLJ22488 fis, clone HRC10	BE672313	2.45

dimethylarginine dimethylaminohydrolase 1	AL078459	2.44
PRO0644 protein	NM_014136	2.44
coat protein gamma-cop	NM_016128	2.43
Consensus includes gb:BG403790 /FEA=EST /DB_XREF=gi:13297238 /DB_XREF=est:602419627F1 /CLONE=IMAGE:4526599 /UG=Hs.158154 ESTs	BG403790	2.43
sema domain, immunoglobulin domain (Ig), short basic domain, secreted, (semaphorin) 3C	NM_006379	2.43
UDP-Gal:betaGlcNAc beta 1,3-galactosyltransferase, polypeptide 5	NM_006057	2.42
hypothetical protein FLJ20059	NM_017644	2.42
CDC14 cell division cycle 14 homolog A (S. cerevisiae)	NM_003672	2.41
hypothetical protein FLJ23548	NM_024590	2.41
hypothetical protein FLJ22558	NM_022747	2.41
nuclear factor I/B	U70862	2.40
hypothetical protein MyO14	NM_030918	2.40
hypothetical protein FLJ20006	NM_017618	2.40
WNT1 inducible signaling pathway protein 3	AF143679	2.40
dickkopf homolog 2 (Xenopus laevis)	NM_014421	2.39
Consensus includes gb:AK021440.1 /DEF=Homo sapiens cDNA FLJ11378 fis, clone HEMBA1000456. /FEA=mRNA /DB_XREF=gi:10432625 /UG=Hs.6937 hypothetical protein FLJ10276	AK021440	2.39
Consensus includes gb:AL359578.1 /DEF=Homo sapiens mRNA; cDNA DKFZp547N163 (from clone DKFZp547N163). /FEA=mRNA /DB_XREF=gi:8655637 /UG=Hs.306511 Homo sapiens mI	AL359578	2.39
Consensus includes gb:AA629050 /FEA=EST /DB_XREF=gi:2541437 /DB_XREF=est:zu84a06.s1 /CLONE=IMAGE:744658 /UG=Hs.50760 ESTs, Highly similar to BimL H.sapiens	AA629050	2.39
Dystonia musculorum of mouse, human homolog of	AL049215	2.39
Consensus includes gb:AF131777.1 /DEF=Homo sapiens clone 25061 mRNA sequence /FEA=mRNA /DB_XREF=gi:4406602 /UG=Hs.183475 Homo sapiens clone 25061 mRNA sequence	AF131777	2.39
hypothetical protein FLJ22746	NM_024785	2.38
H1 histone family, member 4	AL353759	2.38
unc-51-like kinase 2 (C. elegans)	BG526973	2.38
zinc finger protein 42 (myeloid-specific retinoic acid- responsive)	AI733248	2.38
checkpoint suppressor 1	AA860806	2.37
protease, serine, 21 (testisin)	NM_006799	2.37
peptidyl arginine deiminase, type V	AF229067	2.37
calpain 9 (nCL-4)	AB038463	2.37
sulfotransferase family, cytosolic, 1A, phenol-preferring, member 3	L25275	2.37
Eph83	X75208	2.37
craniofacial development protein 1	D85939	2.36
sialophorin (gpL115, leukosialin, CD43)	X60502	2.36
Consensus includes gb:K00627.1 /DEF=human kpni repeat mrna (cdna clone pcd-kpni-8), 3 end. /FEA=mRNA /DB_XREF=gi:337653 /UG=Hs.203776 Human kpni repeat mrna (cdna clone)	K00627	2.36
Consensus includes gb:AK022045.1 /DEF=Homo sapiens cDNA FLJ11983 fis, clone HEMBB1001337. /FEA=mRNA /DB_XREF=gi:10433364 /UG=Hs.193162 Homo sapiens cDNA FLJ11983 f	AK022045	2.36
hypothetical protein FLJ20097	NM_017667	2.36
Consensus includes gb:M78162 /FEA=EST /DB_XREF=gi:273899 /DB_XREF=est:EST01755 /CLONE=HHCPN60 /UG=Hs.314534 ESTs, Moderately similar to ALU5_HUMAN ALU SUBFAMILY S	M78162	2.35
hypothetical protein FLJ20019	NM_017624	2.35
calcium/calmodulin-dependent serine protein kinase (MAGUK family)	AB039327	2.35
M10098 Human 18S rRNA gene, complete (_5, _M, _3 represent transcript regions 5 prime, Middle, and 3 prime respectively)	M10098	2.35
Consensus includes gb:S80491.1 /DEF=stem cell factor (alternatively spliced) human, preimplantation embryos, blastocysts, mRNA Partial, 180 nt. /FEA=mRNA /GEN=stem cell factor, SCF .	S80491	2.35
hemoglobin, gamma G	NM_000184	2.34
transcription factor 20 (AR1)	U19345	2.34
Consensus includes gb:AK023783.1 /DEF=Homo sapiens cDNA FLJ13721 fis, clone PLACE2000450. /FEA=mRNA /DB_XREF=gi:10435820 /UG=Hs.289035 Homo sapiens cDNA FLJ13721 fis	AK023783	2.34
CGI-58 protein	NM_016006	2.34
hypothetical protein FLJ10254	NM_018041	2.34
interleukin 1 receptor antagonist	BE563442	2.33
Bloom syndrome	NM_000057	2.33
Consensus includes gb:AA780524 /FEA=EST /DB_XREF=gi:2839855 /DB_XREF=est:ac71f01.s1 /CLONE=IMAGE:868057 /UG=Hs.294072 ESTs, Weakly similar to ALU1_HUMAN ALU SUBFAI	AA780524	2.33
potassium inwardly-rectifying channel, subfamily J, member 8	NM_004982	2.33
hypothetical protein FLJ14310	NM_025028	2.33
thrombospondin 1	NM_003246	2.33
calmodulin 1 (phosphorylase kinase, delta)	M27319	2.32
leukocyte immunoglobulin-like receptor, subfamily B (with TM and ITIM domains), member 3	AF009643	2.32
M10098 Human 18S rRNA sequence, length 1969 bases, 3 prime target bases 1293-1938	M10098	2.32
hypothetical protein FLJ23447	NM_024825	2.32
KIAA0304 gene product	AF105279	2.32
a disintegrin and metalloproteinase domain 17 (tumor necrosis factor, alpha, converting enzyme)	NM_003183	2.32
SRY (sex determining region Y)-box 21	NM_007084	2.32
phosphatidylinositol-4-phosphate 5-kinase, type II, alpha	NM_005028	2.32
homeo box C8	NM_022658	2.32
CGI-86 protein	BC000637	2.32
guanine nucleotide binding protein 13, gamma	NM_016541	2.31
potassium voltage-gated channel, delayed-rectifier, subfamily S, member 3	NM_002252	2.31

Consensus includes gb:AK000864.1 /DEF=Homo sapiens cDNA FLJ10002 fis, clone HEMBA1000046. /FEA=mRNA /DB_XREF=gi:7021188 /UG=Hs.296522 Homo sapiens cDNA FLJ10002 fis	AK000864	2.31
nuclear LIM interactor-interacting factor	AF229163	2.31
Consensus includes gb:AW451711 /FEA=EST /DB_XREF=gi:6992487 /DB_XREF=est:UI-H-BI3-ale-c-02-0-UI.s1 /CLONE=IMAGE:2736386 /UG=Hs.313760 ESTs	AW451711	2.31
Consensus includes gb:AW090043 /FEA=EST /DB_XREF=gi:6047387 /DB_XREF=est:xd01c05.x1 /CLONE=IMAGE:2592488 /UG=Hs.326464 Homo sapiens cDNA: FLJ21702 fis, clone COL09	AW090043	2.31
elaC homolog 1 (E. coli)	NM_018696	2.31
coagulation factor II (thrombin) receptor	NM_001992	2.30
Consensus includes gb:AK001448.1 /DEF=Homo sapiens cDNA FLJ10586 fis, clone NT2RP2003986. /FEA=mRNA /DB_XREF=gi:7022713 /UG=Hs.301255 Homo sapiens cDNA FLJ10586 fis	AK001448	2.30
gb:NM_018541.1 /DEF=Homo sapiens hypothetical protein PRO2832 (PRO2832), mRNA. /FEA=mRNA /GEN=PRO2832 /PROD=hypothetical protein PRO2832 /DB_XREF=gi:8924207 /UG=	NM_018541	2.30
artemis protein	AK022922	2.30
G protein-coupled receptor 1	AL046992	2.30
KIAA0189 gene product	NM_014725	2.30
protein phosphatase, EF hand calcium-binding domain 2	NM_006239	2.29
nudix (nucleoside diphosphate linked moiety X)-type motif 4	AB007956	2.29
collectin sub-family member 12	NM_030781	2.29
synaptojanin 2	AK026758	2.29
ficolin (collagen/fibrinogen domain-containing lectin) 2 (hucolin)	NM_004108	2.29
KIAA0328 protein	AB002326	2.28
Consensus includes gb:U80770.1 /DEF=Human EST clone 251800 mariner transposon Hsmar1 sequence. /FEA=mRNA /DB_XREF=gi:2921608 /UG=Hs.42192 Human EST clone 251800 ma	U80770	2.28
choline/ethanolaminephosphotransferase	NM_006090	2.28
dopamine receptor D4	NM_000797	2.28
cholinergic receptor, nicotinic, alpha polypeptide 3	M37981	2.28
TXK tyrosine kinase	AW276646	2.28
v-myc myelocytomatosis viral oncogene homolog 2 (avian)	NM_005377	2.27
KIAA0997 protein	NM_014950	2.27
glutamate receptor, metabotropic 8	AI236922	2.27
hypothetical protein FLJ13769	NM_025012	2.27
RAD54, S. cerevisiae, homolog of, B	NM_012415	2.27
GDNF family receptor alpha 1	NM_005264	2.26
Consensus includes gb:AI393960 /FEA=EST /DB_XREF=gi:4223507 /DB_XREF=est:tg11d04.x1 /CLONE=IMAGE:2108455 /UG=Hs.274851 ESTs	AI393960	2.26
Consensus includes gb:AW970881 /FEA=EST /DB_XREF=gi:8160726 /DB_XREF=est:EST382964 /UG=Hs.205660 ESTs	AW970881	2.26
Consensus includes gb:AL137590.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434K0610 (from clone DKFZp434K0610). /FEA=mRNA /DB_XREF=gi:6808330 /UG=Hs.300785 Homo sapiens	AL137590	2.26
lymphocyte cytosolic protein 2 (SH2 domain-containing leukocyte protein of 76kD)	AI123251	2.26
signal transduction protein (SH3 containing)	AB001467	2.25
adenomatosis polyposis coli	S67788	2.25
PRO0097 protein	NM_014114	2.25
RAB36, member RAS oncogene family	AF133588	2.25
transmembrane activator and CAML interactor	NM_012452	2.25
Consensus includes gb:AV702306 /FEA=EST /DB_XREF=gi:10718636 /DB_XREF=est:AV702306 /CLONE=ADBBIDO5 /UG=Hs.321211 ESTs, Highly similar to zinc finger protein 216 splice v	AV702306	2.25
H3 histone family, member C	NM_003531	2.24
hypothetical protein FLJ12975	NM_024809	2.24
hypothetical protein FLJ14346	NM_025029	2.24
Consensus includes gb:AL117451.1 /DEF=Homo sapiens mRNA; cDNA DKFZp586E2317 (from clone DKFZp586E2317). /FEA=mRNA /DB_XREF=gi:5911901 /UG=Hs.293563 Homo sapiens	AL117451	2.24
peptide YY, 2 (seminalplasmin)	NM_021093	2.24
CD34 antigen	M81104	2.24
matrix metalloproteinase 25	NM_022718	2.24
kinectin 1 (kinesin receptor)	BF589024	2.24
NG37 protein	NM_025258	2.24
neuronal cell adhesion molecule	NM_005010	2.24
PAX transcription activation domain interacting protein 1 like	AI357401	2.24
hypothetical protein FLJ13162	NM_025002	2.23
Consensus includes gb:AW173720 /FEA=EST /DB_XREF=gi:6439668 /DB_XREF=est:xj11f05.x1 /CLONE=IMAGE:2656929 /UG=Hs.271356 ESTs	AW173720	2.23
RNA-binding protein gene with multiple splicing	D84109	2.23
Consensus includes gb:AV726900 /FEA=EST /DB_XREF=gi:10836321 /DB_XREF=est:AV726900 /CLONE=HTCABD05 /UG=Hs.7000 ESTs	AV726900	2.23
Consensus includes gb:AI686582 /FEA=EST /DB_XREF=gi:4897876 /DB_XREF=est:tr08e05.x1 /CLONE=IMAGE:2268608 /UG=Hs.283824 Homo sapiens Skn-1ai-like mRNA	AI686582	2.23
distal-less homeo box 2	NM_004405	2.23
T cell activation, increased late expression	NM_005816	2.23
Consensus includes gb:AV708130 /FEA=EST /DB_XREF=gi:10725395 /DB_XREF=est:AV708130 /CLONE=ADCAMA05 /UG=Hs.283965 Homo sapiens scavenger receptor class B type III SR-I	AV708130	2.23
discs, large (Drosophila) homolog-associated protein 1	NM_004746	2.23
leucine zipper-like protein	NM_013344	2.23
Consensus includes gb:AU145680 /FEA=EST /DB_XREF=gi:11007201 /DB_XREF=est:AU145680 /CLONE=HEMBA1005500 /UG=Hs.306613 Homo sapiens cDNA FLJ11740 fis, clone HEMBA	AU145680	2.23
chymotrypsin C (caldecrin)	NM_007272	2.23
golgi associated, gamma adaptin ear containing, ARF binding protein 2	AI799984	2.22
hypothetical protein FLJ12619	AL136632	2.22

zinc finger protein 175	NM_007147	2.22
Consensus includes gb:AK025360.1 /DEF=Homo sapiens cDNA: FLJ21707 fis, clone COL09953. /FEA=mRNA /DB_XREF=gi:10437861 /UG=Hs.306806 Homo sapiens cDNA: FLJ21707 fis, cl	AK025360	2.22
UDP glycosyltransferase 2 family, polypeptide B17	NM_001077	2.22
SRY (sex determining region Y)-box 4	AI989477	2.22
cyclin-dependent kinase inhibitor 2A (melanoma, p16, inhibits CDK4)	AF115544	2.21
phosphoenolpyruvate carboxykinase 1 (soluble)	NM_002591	2.21
RAD1 homolog (S. pombe)	NM_002853	2.21
KIAA0416 protein	NM_015564	2.21
putative zinc finger protein from EUROIMAGE 566589	NM_021220	2.21
3'(2'), 5'-bisphosphate nucleotidase 1	NM_006085	2.21
AT2 receptor-interacting protein 1	BE552421	2.21
makorin, ring finger protein, 4	NM_030757	2.21
hypothetical protein FLJ22479	NM_024900	2.21
melanocortin 1 receptor (alpha melanocyte stimulating hormone receptor)	BG034972	2.20
Consensus includes gb:AC004460 /DEF=Homo sapiens PAC clone RP5-1086D14 /FEA=CDS /DB_XREF=gi:2981263 /UG=Hs.307352 Homo sapiens PAC clone RP5-1086D14	AC004460	2.20
gamma-glutamyltransferase 2	AI344075	2.20
guanine nucleotide binding protein 4	NM_004485	2.20
Parkinson disease (autosomal recessive, juvenile) 2, parkin	NM_004562	2.20
glycoprotein M6B	AI419030	2.20
carboxypeptidase M	NM_001874	2.20
Consensus includes gb:AU148005 /FEA=EST /DB_XREF=gi:11009526 /DB_XREF=est:AU148005 /CLONE=MAMMA1002355 /UG=Hs.296718 Homo sapiens cDNA FLJ12359 fis, clone MAMM	AU148005	2.20
Consensus includes gb:AF052108.1 /DEF=Homo sapiens clone 23687 mRNA sequence. /FEA=mRNA /DB_XREF=gi:3360415 /UG=Hs.12513 Homo sapiens clone 23687 mRNA sequence	AF052108	2.20
DNA fragmentation factor, 45 kD, alpha polypeptide	NM_004401	2.20
receptor tyrosine kinase-like orphan receptor 1	BC006374	2.19
Consensus includes gb:N46852 /FEA=EST /DB_XREF=gi:1188018 /DB_XREF=est:yy73d05.s1 /CLONE=IMAGE:279177 /UG=Hs.285625 Homo sapiens mRNA; cDNA DKFZp434A119 (from cl	N46852	2.19
SCAN domain-containing 2	NM_022050	2.19
KIAA1028 protein	AY028424	2.19
hypothetical protein R32184_1	AC004528	2.19
HYA22 protein	BF002474	2.19
hypothetical protein PRO2900	NM_018635	2.19
chromosome 21 open reading frame 62	NM_019596	2.19
calcitonin receptor	NM_001742	2.19
carcinoembryonic antigen-related cell adhesion molecule 1 (biliary glycoprotein)	D12502	2.19
Alu-binding protein with zinc finger domain	NM_014274	2.18
neural cell adhesion molecule 2	NM_004540	2.18
glutathione peroxidase 2 (gastrointestinal)	NM_002083	2.18
KIAA0870 protein	AI808958	2.18
intracellular hyaluronan-binding protein	AF241831	2.18
matrix metalloproteinase 10 (stromelysin 2)	NM_002425	2.18
bridging integrator 2	NM_016293	2.18
glutamate receptor, ionotropic, kainate 4	NM_014619	2.18
antigen identified by monoclonal antibody MRC OX-2	H23979	2.18
Consensus includes gb:AL512703.1 /DEF=Homo sapiens mRNA; cDNA DKFZp667H046 (from clone DKFZp667H046). /FEA=mRNA /GEN=DKFZp667H046 /PROD=hypothetical protein /DB_X	AL512703	2.18
M11167 Human 28S rRNA sequence, length 5025 bases, 3 prime target bases 3331-4995	M11167	2.18
solute carrier family 14 (urea transporter), member 1 (Kidd blood group)	NM_015865	2.17
Consensus includes gb:AK025072.1 /DEF=Homo sapiens cDNA: FLJ21419 fis, clone COL04084. /FEA=mRNA /DB_XREF=gi:10437512 /UG=Hs.306759 Homo sapiens cDNA: FLJ21419 fis, cl	AK025072	2.17
Consensus includes gb:AK001442.1 /DEF=Homo sapiens cDNA FLJ10580 fis, clone NT2RP2003533. /FEA=mRNA /DB_XREF=gi:7022701 /UG=Hs.140402 Homo sapiens cDNA FLJ10580 fis	AK001442	2.17
KIAA0341 protein	AW139448	2.17
solute carrier family 9 (sodium/hydrogen exchanger), isoform 2	AF073299	2.17
gb:NM_017648.1 /DEF=Homo sapiens hypothetical protein FLJ20063 (FLJ20063), mRNA. /FEA=mRNA /GEN=FLJ20063 /PROD=hypothetical protein FLJ20063 /DB_XREF=gi:8923068 /UG=	NM_017648	2.17
gb:U96291.1 /DEF=Homo sapiens Ig kappa light chain variable region (VkII-A23) mRNA, complete cds. /FEA=mRNA /GEN=VkII-A23 /PROD=Ig kappa light chain variable region /DB_XREF=.	U96291	2.17
Consensus includes gb:AL136105 /DEF=Human DNA sequence from clone RP4-670F13 on chromosome 1q42.2-43. Contains an enolase 1, (alpha) (ENO1) pseudogene, the gene for Po66	AL136105	2.16
golgi phosphoprotein 4	NM_014498	2.16
Consensus includes gb:AK022219.1 /DEF=Homo sapiens cDNA FLJ12157 fis, clone MAMMA1000500. /FEA=mRNA /DB_XREF=gi:10433569 /UG=Hs.98812 Homo sapiens cDNA FLJ12157 fis	AK022219	2.16
interleukin 1, beta	M15330	2.16
Consensus includes gb:AK025077.1 /DEF=Homo sapiens cDNA: FLJ21424 fis, clone COL04157. /FEA=mRNA /DB_XREF=gi:10437519 /UG=Hs.287662 Homo sapiens cDNA: FLJ21424 fis, cl	AK025077	2.16
Consensus includes gb:AK025038.1 /DEF=Homo sapiens cDNA: FLJ21385 fis, clone COL03356. /FEA=mRNA /DB_XREF=gi:10437474 /UG=Hs.306750 Homo sapiens cDNA: FLJ21385 fis, cl	AK025038	2.15
mitogen-activated protein kinase 8 interacting protein 2	NM_016431	2.15
Consensus includes gb:BF693921 /FEA=EST /DB_XREF=gi:11979329 /DB_XREF=est:602082488F1 /CLONE=IMAGE:4246917 /UG=Hs.180513 ATP-binding cassette, sub-family A (ABC1),	BF693921	2.15
M27830 Human 28S ribosomal RNA gene, complete cds (_5, _M, _3 represent transcript regions 5 prime, Middle, and 3 prime respectively)	M27830	2.15
hematopoietic PBX-interacting protein	AI348545	2.15
potassium inwardly-rectifying channel, subfamily J, member 5	NM_000890	2.15
solute carrier family 1 (glial high affinity glutamate transporter), member 2	NM_004171	2.15

serine/threonine kinase 6	NM_003158	2.15
v-raf-1 murine leukemia viral oncogene homolog 1	AI088622	2.15
UDP-Gal:betaGlcNAc beta 1,3-galactosyltransferase, polypeptide 3	AB050855	2.15
Consensus includes gb:AV756729 /FEA=EST /DB_XREF=gi:10914577 /DB_XREF=est:AV756729 /CLONE=BMCACB11 /UG=Hs.317661 ESTs	AV756729	2.15
SH3 adaptor protein (a Src homology 2 protein)	NM_003028	2.14
pleiomorphic adenoma gene 1	NM_002655	2.14
manic fringe homolog (Drosophila)	AI760053	2.14
PDZ domain containing guanine nucleotide exchange factor(GEF)1	AL117397	2.14
Consensus includes gb:AW971254 /FEA=EST /DB_XREF=gi:8161099 /DB_XREF=est:EST383343 /UG=Hs.178433 ESTs	AW971254	2.14
Consensus includes gb:AL050154.1 /DEF=Homo sapiens mRNA; cDNA DKFZp586L0120 (from clone DKFZp586L0120). /FEA=mRNA /DB_XREF=gi:4884366 /UG=Hs.98314 Homo sapiens m	AL050154	2.14
proprotein convertase subtilisin/kexin type 1	NM_000439	2.14
hydroxy-delta-5-steroid dehydrogenase, 3 beta- and steroid delta-isomerase 2	NM_000198	2.14
aldehyde dehydrogenase 3 family, member B1	BC002553	2.14
KIAA0460 protein	AK026120	2.14
interleukin 4	NM_000589	2.14
phosducin-like	NM_005388	2.14
plasminogen activator, urokinase	NM_002658	2.14
apolipoprotein F	NM_001638	2.14
general transcription factor IIH, polypeptide 3 (34kD subunit)	AI569458	2.13
neuregulin 2	NM_013984	2.13
cell cycle progression 8 protein	NM_004748	2.13
Consensus includes gb:X07882 /DEF=Human PRB4 gene for proline-rich protein Po, allele 5 /FEA=mRNA /DB_XREF=gi:35647 /UG=Hs.103972 proline-rich protein BstNI subfamily 4	X07882	2.13
Consensus includes gb:AK023017.1 /DEF=Homo sapiens cDNA FLJ12955 fis, clone NT2RP2005496, moderately similar to ZINC FINGER PROTEIN 135. /FEA=mRNA /DB_XREF=gi:10434737	AK023017	2.13
KIAA0906 protein	AA909765	2.13
hypothetical protein FLJ11155	NM_018342	2.13
hypothetical protein FLJ21596	NM_024823	2.13
zinc finger protein 236	AK000847	2.13
Consensus includes gb:AU146646 /FEA=EST /DB_XREF=gi:11008167 /DB_XREF=est:AU146646 /CLONE=HEMBB1001096 /UG=Hs.179752 Homo sapiens cDNA FLJ10270 fis, clone HEMBB	AU146646	2.13
IGF-II mRNA-binding protein 2	NM_006548	2.13
neurotrophic tyrosine kinase, receptor, type 3	NM_002530	2.13
KIAA0610 protein	AK002207	2.13
transcription factor (p38 interacting protein)	NM_017569	2.13
ectodermal dysplasia 1, anhidrotic	NM_001399	2.13
hypothetical protein	NM_020153	2.13
oligophrenin 1	NM_002547	2.12
calcium channel, voltage-dependent, beta 4 subunit	NM_000726	2.12
HCGII-7 protein	X81001	2.12
mannosidase, alpha, class 1A, member 2	AI189305	2.12
hypothetical protein FLJ20274	NM_017736	2.12
DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 34	NM_014681	2.11
taste receptor, type 2, member 16	NM_016945	2.11
zinc finger protein 223	NM_013361	2.11
brain-specific angiogenesis inhibitor 3	AB011122	2.11
Consensus includes gb:AI042030 /FEA=EST /DB_XREF=gi:3281224 /DB_XREF=est:oz60a09.x1 /CLONE=IMAGE:1660696 /UG=Hs.132153 ESTs	AI042030	2.11
DKFZp434P211 protein	AL117484	2.11
Williams-Beuren syndrome chromosome region 23	NM_025042	2.11
Consensus includes gb:AK025095.1 /DEF=Homo sapiens cDNA: FLJ21442 fis, clone COL04429, highly similar to HSA237839 Homo sapiens mRNA for hypothetical protein. /FEA=mRNA /DB	AK025095	2.11
patched homolog (Drosophila)	AL044175	2.11
Consensus includes gb:AF070547.1 /DEF=Homo sapiens clone 24820 mRNA sequence. /FEA=mRNA /DB_XREF=gi:3387909 /UG=Hs.146312 Homo sapiens clone 24820 mRNA sequence	AF070547	2.11
KIAA0429 gene product	AF116674	2.11
DKFZP434C091 protein	AL080170	2.11
sirtuin silent mating type information regulation 2 homolog 4 (S. cerevisiae)	NM_012240	2.11
transcription factor (p38 interacting protein)	AW003334	2.11
Consensus includes gb:AI246331 /FEA=EST /DB_XREF=gi:3841728 /DB_XREF=est:qu39f10.x1 /CLONE=IMAGE:1967179 /UG=Hs.248999 ESTs	AI246331	2.11
RNA binding motif, single stranded interacting protein 2	NM_002898	2.11
hypothetical protein DKFZp434C135	NM_030936	2.11
actin filament associated protein	NM_021638	2.11
tankyrase, TRF1-interacting ankyrin-related ADP-ribose polymerase 2	NM_025235	2.10
hypothetical protein PRO2533	NM_018629	2.10
KIAA0254 gene product	BG434168	2.10
solute carrier family 16 (monocarboxylic acid transporters), member 1	BF511091	2.10
gb:BC000108.1 /DEF=Homo sapiens, Similar to Nedd-4-like ubiquitin-protein ligase, clone MGC:2079, mRNA, complete cds. /FEA=mRNA /PROD=Similar to Nedd-4-like ubiquitin-protein liga	BC000108	2.10
pregnancy specific beta-1-glycoprotein 3	R32065	2.10

hypothetical protein MGC3196	NM_024084	2.10
HCNP protein; XPA-binding protein 2	NM_020196	2.10
KIAA0557 protein	AI871396	2.10
KIAA0007 protein	D26488	2.10
Consensus includes gb:BF035279 /FEA=EST /DB_XREF=gi:10743006 /DB_XREF=est:601457165F1 /CLONE=IMAGE:3860633 /UG=Hs.20325 ESTs, Moderately similar to pot. ORF V H.sapi	BF035279	2.10
Wilms tumor 1	S75264	2.10
CDC14 cell division cycle 14 homolog B (S. cerevisiae)	BF792631	2.10
interleukin 13 receptor, alpha 1	U62858	2.10
homeo box C5	NM_018953	2.10
dipeptidylpeptidase IV (CD26, adenosine deaminase complexing protein 2)	M80536	2.10
basic helix-loop-helix domain containing, class B, 3	BE857425	2.10
Treacher Collins-Franceschetti syndrome 1	AW167713	2.10
aquaporin 4	D63412	2.10
prothymosin a14	NM_016171	2.09
N-myc downstream-regulated gene 2	W74452	2.09
hypothetical protein PRO1584	NM_018586	2.09
ATPase, H+ transporting, lysosomal (vacuolar proton pump) 9kD	AI252582	2.09
Consensus includes gb:AV753204 /FEA=EST /DB_XREF=gi:10911052 /DB_XREF=est:AV753204 /CLONE=NPOBDB08 /UG=Hs.170267 ESTs	AV753204	2.09
Consensus includes gb:AF052160.1 /DEF=Homo sapiens clone 24629 mRNA sequence. /FEA=mRNA /DB_XREF=gi:3360471 /UG=Hs.142570 Homo sapiens clone 24629 mRNA sequence	AF052160	2.09
apolipoprotein C-I	W79394	2.09
H2A histone family, member X	H51429	2.09
gb:BC003528.1 /DEF=Homo sapiens, clone MGC:10782, mRNA, complete cds. /FEA=mRNA /PROD=Unknown (protein for MGC:10782) /DB_XREF=gi:13097626 /UG=Hs.306969 Homo sapi	BC003528	2.09
endothelin converting enzyme 1	NM_001397	2.09
Microfibril-associated glycoprotein-2	AW665892	2.09
fibroblast growth factor 17	NM_003867	2.09
Consensus includes gb:AK022192.1 /DEF=Homo sapiens cDNA FLJ12130 fis, clone MAMMA1000251. /FEA=mRNA /DB_XREF=gi:10433534 /UG=Hs.319567 Homo sapiens cDNA FLJ12130	AK022192	2.09
hypothetical protein FLJ22670	NM_025144	2.08
hypothetical protein LOC57187	BG403671	2.08
Consensus includes gb:AA017093 /FEA=EST /DB_XREF=gi:1479257 /DB_XREF=est:ze37h10.s1 /CLONE=IMAGE:361219 /UG=Hs.271780 ESTs	AA017093	2.08
T-box 5	U80987	2.08
connective tissue growth factor	M92934	2.08
KIAA0319 gene product	NM_014809	2.08
xylulokinase (H. influenzae) homolog	AA777793	2.08
cullin 5	NM_003478	2.08
Consensus includes gb:AK001065.1 /DEF=Homo sapiens cDNA FLJ10203 fis, clone HEMBA1004930, moderately similar to 26S PROTEASOME SUBUNIT S5B. /FEA=mRNA /DB_XREF=gi:7022	AK001065	2.08
vav 2 oncogene	NM_003371	2.08
Consensus includes gb:AV684859 /FEA=EST /DB_XREF=gi:10286722 /DB_XREF=est:AV684859 /CLONE=GKCDIA03 /UG=Hs.251334 ESTs, Moderately similar to ALU6_HUMAN ALU SUBFAM	AV684859	2.08
trichorhinophalangeal syndrome I	NM_014112	2.08
Consensus includes gb:AK025552.1 /DEF=Homo sapiens cDNA: FLJ21899 fis, clone HEP03467. /FEA=mRNA /DB_XREF=gi:10438101 /UG=Hs.306815 Homo sapiens cDNA: FLJ21899 fis, cl	AK025552	2.08
Consensus includes gb:BG433489 /FEA=EST /DB_XREF=gi:13339995 /DB_XREF=est:602497919F1 /CLONE=IMAGE:4611460 /UG=Hs.297257 ESTs	BG433489	2.07
Consensus includes gb:AL050204.1 /DEF=Homo sapiens mRNA; cDNA DKFZp586F1223 (from clone DKFZp586F1223). /FEA=mRNA /DB_XREF=gi:4884443 /UG=Hs.28540 Homo sapiens m	AL050204	2.07
transmembrane protein 8 (five membrane-spanning domains)	AI636233	2.07
PRO1621 protein	AF113018	2.07
dual oxidase 1	AL137592	2.07
CGI-142	BG285017	2.07
caveolin 2	AA150110	2.07
dopamine receptor D2	S62137	2.07
KIAA1644 protein	AL047020	2.07
Consensus includes gb:AK022892.1 /DEF=Homo sapiens cDNA FLJ12830 fis, clone NT2RP2003073. /FEA=mRNA /DB_XREF=gi:10434549 /UG=Hs.123158 Homo sapiens cDNA FLJ12830 fi	AK022892	2.07
SMA3	NM_006780	2.07
homolog of mouse BMP-2 inducible kinase	AB015331	2.07
aldehyde dehydrogenase 4 family, member A1	U24267	2.07
Consensus includes gb:AF319523.1 /DEF=Homo sapiens RT-LI mRNA, complete sequence. /FEA=mRNA /PROD=RT-LI /DB_XREF=gi:13310144 /UG=Hs.326775 Homo sapiens RT-LI mRNA,	AF319523	2.07
core-binding factor, runt domain, alpha subunit 2; translocated to, 1; cyclin D-related	NM_004349	2.07
alpha-2-glycoprotein 1, zinc	D90427	2.07
ubiquitin protein ligase E3A (human papilloma virus E6-associated protein, Angelman syndrome)	BF588511	2.06
phospholipid scramblase 2	NM_020359	2.06
DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 18 (Myc-regulated)	BG530850	2.06
Consensus includes gb:AK022023.1 /DEF=Homo sapiens cDNA FLJ11961 fis, clone HEMB81001020, highly similar to Homo sapiens mRNA for KIAA0889 protein. /FEA=mRNA /DB_XREF=gi	AK022023	2.06
MADS box transcription enhancer factor 2, polypeptide D (myocyte enhancer factor 2D)	NM_005920	2.06
pleckstrin homology-like domain, family A, member 1	AI795908	2.06
cytochrome P450 polypeptide 43	AF280113	2.06
Consensus includes gb:AK022174.1 /DEF=Homo sapiens cDNA FLJ12112 fis, clone MAMMA1000043. /FEA=mRNA /DB_XREF=gi:10433510 /UG=Hs.288793 Homo sapiens cDNA FLJ12112	AK022174	2.06

tuftelin-interacting protein	NM_012143	2.06
HMT1 hnRNP methyltransferase-like 1 (S. cerevisiae)	U79286	2.06
Consensus includes gb:AL078636.1 /DEF=Homo sapiens mRNA full length insert cDNA clone EUROIMAGE 117929. /FEA=mRNA /DB_XREF=gi:5051866 /UG=Hs.227171 Homo sapiens mRN	AL078636	2.06
hypothetical protein FLJ10488	BF591163	2.06
Consensus includes gb:U63332.1 /DEF=Human super cysteine rich protein mRNA, partial cds. /FEA=mRNA /PROD=super cysteine rich protein /DB_XREF=gi:1480862 /UG=Hs.169325 Hum	U63332	2.06
gb:U79250.1 /DEF=Human glycerol-3-phosphate dehydrogenase mRNA, complete cds. /FEA=mRNA /PROD=glycerol-3-phosphate dehydrogenase /DB_XREF=gi:1806627 /FL=gb:U79250.1	U79250	2.06
Consensus includes gb:AL049252.1 /DEF=Homo sapiens mRNA; cDNA DKFZp564D193 (from clone DKFZp564D193). /FEA=mRNA /DB_XREF=gi:4499993 /UG=Hs.302048 Homo sapiens mI	AL049252	2.06
Consensus includes gb:AL035067 /DEF=Human DNA sequence from clone 170F5 on chromosome Xq22.3-24. Contains an HMG1 (high-mobility group (nonhistone chromosomal) protein 1)	AL035067	2.05
intermediate filament protein syncoilin	NM_030786	2.05
PC4 and SFRS1 interacting protein 1	AF098482	2.05
glycoprotein IX (platelet)	NM_000174	2.05
KIAA1102 protein	AK027231	2.05
sorting nexin 15	NM_013306	2.05
hypothetical protein FLJ21290	NM_025034	2.05
dentin matrix acidic phosphoprotein	NM_004407	2.05
Consensus includes gb:R78299 /FEA=EST /DB_XREF=gi:853409 /DB_XREF=est:yi81h12.s1 /CLONE=IMAGE:145703 /UG=Hs.287419 Homo sapiens cDNA FLJ11505 fis, clone HEMBA10021	R78299	2.05
sialic acid binding Ig-like lectin 5	NM_003830	2.05
Bacteriophage /REF=X03453 /DEF=Bacteriophage P1 cre recombinase protein corresponding to nucleotides 581-1001 of X03453 /LEN=1058 (-5 and -3 represent transcript regions 5 prin	X03453	2.05
cantaurin, alpha 2	NM_018404	2.04
Consensus includes gb:AW150065 /FEA=EST /DB_XREF=gi:6197971 /DB_XREF=est:zg48a10.x1 /CLONE=IMAGE:2630778 /UG=Hs.271957 ESTs	AW150065	2.04
hypothetical protein FLJ10496	NM_018114	2.04
Consensus includes gb:AF339807.1 /DEF=Homo sapiens clone IMAGE:26186, mRNA sequence. /FEA=mRNA /DB_XREF=gi:13507345 /UG=Hs.137567 Homo sapiens cDNA FLJ11627 fis, clo	AF339807	2.04
adducin 2 (beta)	NM_017488	2.04
enolase 1, (alpha)	U88968	2.04
chromosome 1 open reading frame 6	AK023354	2.04
interferon induced transmembrane protein 1 (9-27)	AA749101	2.04
Consensus includes gb:D25272.1 /DEF=Homo sapiens mRNA, clone:RES4-16. /FEA=mRNA /DB_XREF=gi:464187 /UG=Hs.174225 Homo sapiens mRNA, clone:RES4-16	D25272	2.04
hypothetical protein FLJ10979	NM_018289	2.04
transcription factor 7-like 2 (T-cell specific, HMG-box)	AI703074	2.04
hypothetical protein PRO1438	NM_018600	2.04
chorionic somatomammotropin hormone-like 1	NM_022579	2.04
f-box and leucine-rich repeat protein 11	AK024505	2.04
PRO0365 protein	NM_014126	2.04
hypothetical protein FLJ20284	NM_017744	2.04
sperm adhesion molecule 1 (PH-20 hyaluronidase, zona pellucida binding)	L13779	2.04
N-myc downstream-regulated gene 2	AI358939	2.04
Consensus includes gb:AU146391 /FEA=EST /DB_XREF=gi:11007912 /DB_XREF=est:AU146391 /CLONE=HEMBB1000343 /UG=Hs.209806 Homo sapiens cDNA FLJ11924 fis, clone HEMBB	AU146391	2.04
hypothetical protein FLJ20084	NM_017659	2.03
Consensus includes gb:AK024136.1 /DEF=Homo sapiens cDNA FLJ14074 fis, clone HEMBB1001869. /FEA=mRNA /DB_XREF=gi:10436442 /UG=Hs.141208 Homo sapiens cDNA FLJ14074 f	AK024136	2.03
cytochrome b-245, beta polypeptide (chronic granulomatous disease)	S67289	2.03
Consensus includes gb:AW974910 /FEA=EST /DB_XREF=gi:8166113 /DB_XREF=est:EST387015 /UG=Hs.298138 ESTs	AW974910	2.03
immunoglobulin heavy constant gamma 3 (G3m marker)	BF002659	2.03
hypothetical protein FLJ23189	NM_025057	2.03
testis specific ankyrin-like protein 1	AL136618	2.03
cyclic nucleotide gated channel alpha 3	NM_001298	2.03
hypothetical protein FLJ11588	NM_024603	2.03
glycerol-3-phosphate dehydrogenase 1 (soluble)	AI368018	2.03
translation initiation factor IF2	BE138647	2.03
Consensus includes gb:BF447983 /FEA=EST /DB_XREF=gi:11513212 /DB_XREF=est:hr83f09.x1 /CLONE=IMAGE:3135113 /UG=Hs.58993 Homo sapiens mRNA sequence, IMAGE clone 417	BF447983	2.03
sema domain, immunoglobulin domain (Ig), short basic domain, secreted, (semaphorin) 3F	NM_004186	2.03
small inducible cytokine subfamily A (Cys-Cys), member 16	NM_004590	2.03
hepatocyte nuclear factor 4, alpha	Z49825	2.03
Consensus includes gb:AL390026 /DEF=Human DNA sequence from clone RP3-336K20 on chromosome 6 Contains parts of 2 genes for novel proteins, ESTs, STSs and GSSs /FEA=CDS /DB_	AL390026	2.03
phospholamban	M60411	2.02
hypothetical protein FLJ23516	NM_024539	2.02
N-acylsphingosine amidohydrolase (acid ceramidase)	U47674	2.02
myosin, heavy polypeptide 13, skeletal muscle	NM_003802	2.02
DKFZP547E1010 protein	N92920	2.02
Consensus includes gb:AL109696.1 /DEF=Homo sapiens mRNA full length insert cDNA clone EUROIMAGE 21920. /FEA=mRNA /DB_XREF=gi:5689771 /UG=Hs.185701 Homo sapiens mRNA	AL109696	2.02
retinoic acid receptor, beta	X04014	2.02
LIM and senescent cell antigen-like domains 1	NM_004987	2.02
protein kinase, cGMP-dependent, type I	NM_006258	2.02
KIAA1016 protein	AW298219	2.02

Consensus includes gb:AL390857 /DEF=Human DNA sequence from clone RP11-51N22 on chromosome 13 Contains ESTs, STSs and GSSs. Contains an HNRPA1 (heterogeneous nuclear ribo	AL390857	2.02
midline 2	AL034399	2.02
Consensus includes gb:AL359563.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434D0326 (from clone DKFZp434D0326). /FEA=mRNA /DB_XREF=gi:8655619 /UG=Hs.306509 Homo sapiens	AL359563	2.02
SPARC-like 1 (mast9, hevin)	NM_004684	2.02
hypothetical protein FLJ10697	NM_018181	2.02
myristoylated alanine-rich protein kinase C substrate	AA770596	2.02
KIAA0579 protein	AL117532	2.02
sodium bicarbonate transporter 4	AF243499	2.02
Consensus includes gb:AK000861.1 /DEF=Homo sapiens cDNA FLJ20854 fis, clone ADKA01341. /FEA=mRNA /DB_XREF=gi:7021185 /UG=Hs.306412 Homo sapiens cDNA FLJ20854 fis, clo	AK000861	2.02
sterile-alpha motif and leucine zipper containing kinase AZK	NM_016653	2.01
apolipoprotein B mRNA editing enzyme, catalytic polypeptide 1	NM_001644	2.01
TGFB-induced factor (TALE family homeobox)	NM_003244	2.01
hypothetical protein MGC5384	NM_030972	2.01
tripartite motif-containing 14	AF220130	2.01
hypothetical protein	AL079292	2.01
Consensus includes gb:AL049435.1 /DEF=Homo sapiens mRNA; cDNA DKFZp586B0220 (from clone DKFZp586B0220). /FEA=mRNA /DB_XREF=gi:4500218 /UG=Hs.170056 Homo sapiens	AL049435	2.01
glutamate receptor, metabotropic 1	NM_000838	2.01
heparan sulfate (glucosamine) 3-O-sulfotransferase 1	NM_005114	2.01
hypothetical protein FLJ14166	NM_024565	2.01
S100 calcium binding protein, beta (neural)	BC001766	2.01
WW domain binding protein 4 (formin binding protein 21)	AJ734228	2.01
Consensus includes gb:AL359578.1 /DEF=Homo sapiens mRNA; cDNA DKFZp547N163 (from clone DKFZp547N163). /FEA=mRNA /DB_XREF=gi:8655637 /UG=Hs.306511 Homo sapiens mI	AL359578	2.01
neuronal Shc	NM_016848	2.01
Consensus includes gb:AL136094 /DEF=Human DNA sequence from clone RP3-527B10 on chromosome 6q25.1-25.3 Contains a pseudogene similar to HMG (high mobility group) protein, S	AL136094	2.01
myeloid/lymphoid or mixed-lineage leukemia (trithorax homolog, Drosophila); translocated to, 10	AF060938	2.01
KIAA0806 gene product	NM_014813	2.01
nuclear receptor coactivator 1	AJ672428	2.01
KIAA0677 gene product	BC002558	2.00
gb:AF119914.1 /DEF=Homo sapiens PRO3078 mRNA, complete cds. /FEA=mRNA /PROD=PRO3078 /DB_XREF=gi:7770264 /UG=Hs.306573 Homo sapiens PRO3078 mRNA, complete cds /	AF119914	2.00
peter pan homolog (Drosophila)	AB022847	2.00
myelin protein zero-like 1	AF181660	2.00
rho/rac guanine nucleotide exchange factor (GEF) 2	NM_004723	2.00
RNA helicase	NM_014314	2.00
Consensus includes gb:AA053967 /FEA=EST /DB_XREF=gi:1544893 /DB_XREF=est:ze75h04.s1 /CLONE=IMAGE:364855 /UG=Hs.51515 Homo sapiens mRNA; cDNA DKFZp564G112 (from	AA053967	2.00
cystic fibrosis transmembrane conductance regulator, ATP-binding cassette (sub-family C, member 7)	NM_000492	2.00
sialyltransferase 8 (alpha-2, 8-sialytransferase) B	NM_006011	2.00
Bacteriophage /REF=X03453 /DEF=Bacteriophage P1 cre recombinase protein corresponding to nucleotides 1032-1270 of X03453 /LEN=1058 (-5 and -3 represent transcript regions 5 pri	X03453	2.00
hypothetical protein DKFZp761C07121	NM_017594	2.00
Consensus includes gb:AU156204 /FEA=EST /DB_XREF=gi:11017725 /DB_XREF=est:AU156204 /CLONE=PLACE1002583 /UG=Hs.306651 Homo sapiens cDNA FLJ13435 fis, clone PLACE1	AU156204	2.00
desmin	AK022087	2.00
tumor endothelial marker 7 precursor	AF070526	2.00
Consensus includes gb:AK026847.1 /DEF=Homo sapiens cDNA: FLJ23194 fis, clone REC00490. /FEA=mRNA /DB_XREF=gi:10439802 /UG=Hs.306887 Homo sapiens cDNA: FLJ23194 fis, cl	AK026847	2.00
gap junction protein, alpha 1, 43kD (connexin 43)	NM_000165	2.00
KIAA0894 protein	NM_014896	2.00
ATPase, Ca++ transporting, plasma membrane 2	R52647	2.00
chromosome 4 open reading frame 5	AY013295	2.00
hypothetical protein FLJ21075	NM_025031	2.00
short stature homeobox 2	NM_006884	2.00
v-ets erythroblastosis virus E26 oncogene like (avian)	M21535	2.00
Consensus includes gb:U63542.1 /DEF=Human putative FAP protein mRNA, partial cds. /FEA=mRNA /PROD=FAP protein /DB_XREF=gi:1890646 /UG=Hs.166101 Human putative FAP prote	U63542	2.00
Consensus includes gb:AA401963 /FEA=EST /DB_XREF=gi:2055965 /DB_XREF=est:zu53c08.s1 /CLONE=IMAGE:741710 /UG=Hs.194107 ESTs, Moderately similar to ALU7_HUMAN ALU SU	AA401963	2.00
Consensus includes gb:AF257099 /DEF=Homo sapiens prothymosin alpha (PTMA) gene, complete cds /FEA=CDS /DB_XREF=gi:8037944 /UG=Hs.283947 Homo sapiens prothymosin alpha	AF257099	2.00
very low density lipoprotein receptor	L22431	1.99
gb:BC005233.1 /DEF=Homo sapiens, clone MGC:12257, mRNA, complete cds. /FEA=mRNA /PROD=Unknown (protein for MGC:12257) /DB_XREF=gi:13528866 /UG=Hs.326711 Homo sapi	BC005233	1.99
chemokine (C-C motif) receptor 2	NM_000648	1.99
GABA(A) receptor-associated protein like 1	AF087847	1.99
Consensus includes gb:AK025194.1 /DEF=Homo sapiens cDNA: FLJ21541 fis, clone COL06166. /FEA=mRNA /DB_XREF=gi:10437657 /UG=Hs.306784 Homo sapiens cDNA: FLJ21541 fis, cl	AK025194	1.99
dipeptidase 1 (renal)	NM_004413	1.99
Consensus includes gb:BE892698 /FEA=EST /DB_XREF=gi:10353302 /DB_XREF=est:601433917F1 /CLONE=IMAGE:3918989 /UG=Hs.129709 Homo sapiens chromosome 19, cosmid R30	BE892698	1.99
Consensus includes gb:S65921.1 /DEF=anti-colorectal carcinoma light chain=glycoprotein CANAG-50 specific IgG1 kappa human, 19.9 hybridoma, antibody 1116NS19.9, mRNA, 998 nt. /FE	S65921	1.99
zinc finger protein 257	AF070651	1.99
zinc finger protein 272	X78931	1.99
spectrin, beta, erythrocytic (includes spherocytosis, clinical type I)	NM_000347	1.99

par-6 partitioning defective 6 homolog alpha (C.elegans)	NM_016948	1.99
triple functional domain (PTPRF interacting)	AL161955	1.99
Consensus includes gb:AK026913.1 /DEF=Homo sapiens cDNA: FLJ23260 fis, clone COL05804, highly similar to HSU90911 Human clone 23652 mRNA sequence. /FEA=mRNA /DB_XREF=g	AK026913	1.98
v-src sarcoma (Schmidt-Ruppin A-2) viral oncogene homolog (avian)	NM_005417	1.98
potassium voltage-gated channel, Shab-related subfamily, member 1	L02840	1.98
FK506 binding protein 8 (38kD)	N95418	1.98
Consensus includes gb:AL031224 /DEF=Human DNA sequence from clone 336H9 on chromosome 6p12.1-21.1. Contains the alternatively spliced TFAP2B gene for transcription factor AP-2 l	AL031224	1.98
6 protein-coupled receptor 20	NM_005293	1.98
pleckstrin 2 (mouse) homolog	NM_016445	1.98
solute carrier family 10 (sodium/bile acid cotransporter family), member 1	NM_003049	1.98
DKFZP564B0769 protein	AA902326	1.98
Consensus includes gb:AB000464.1 /DEF=Homo sapiens mRNA, exon 1, 2, 3, 4, clone:RES4-24A. /FEA=mRNA /DB_XREF=gi:1843395 /UG=Hs.104258 Homo sapiens mRNA, exon 1, 2, 3, 4	AB000464	1.98
Consensus includes gb:AK026667.1 /DEF=Homo sapiens cDNA: FLJ23014 fis, clone LNG00806. /FEA=mRNA /DB_XREF=gi:10439568 /UG=Hs.306862 Homo sapiens cDNA: FLJ23014 fis, cl	AK026667	1.98
angiopoietin-like 2	NM_012098	1.98
rabaptin-5	NM_004703	1.98
transcription factor 7 (T-cell specific, HMG-box)	NM_003202	1.98
KIAA0669 gene product	NM_014779	1.98
peptidyl-prolyl isomerase 6 (cyclophilin 6)	AW340788	1.98
enhancer of zeste homolog 1 (Drosophila)	AB004818	1.98
Consensus includes gb:AL554245 /FEA=EST /DB_XREF=gi:12894837 /DB_XREF=est:AL554245 /CLONE=CS0DI082YA02 (5 prime) /UG=Hs.288869 nuclear receptor subfamily 2, group F,	AL554245	1.98
periaxin	AB046840	1.98
KIAA0953 protein	AB023170	1.98
period homolog 3 (Drosophila)	Z98884	1.98
carboxypeptidase B1 (tissue)	NM_001871	1.98
KIAA1751 protein	AB051538	1.98
hypothetical protein MGC3101	AK022644	1.98
KIAA1096 protein	AW238632	1.98
chromosome 6 open reading frame 32	AB002384	1.98
flavin containing monooxygenase 2	BC005894	1.98
putative lymphocyte G0/G1 switch gene	NM_015714	1.98
actin related protein 2/3 complex, subunit 1B (41 kD)	AA464753	1.97
gb:BC005907.1 /DEF=Homo sapiens, Similar to histamine N-methyltransferase, clone MGC:14500, mRNA, complete cds. /FEA=mRNA /PROD=Similar to histamine N-methyltransferase /DB_;	BC005907	1.97
glycophorin A (includes MN blood group)	U00179	1.97
hypothetical protein FLJ20700	NM_017932	1.97
Consensus includes gb:AK024561.1 /DEF=Homo sapiens cDNA: FLJ20908 fis, clone ADSE00417. /FEA=mRNA /DB_XREF=gi:10436870 /UG=Hs.306689 Homo sapiens cDNA: FLJ20908 fis,	AK024561	1.97
aldehyde dehydrogenase 1 family, member A2	NM_003888	1.97
Consensus includes gb:AU145135 /FEA=EST /DB_XREF=gi:11006656 /DB_XREF=est:AU145135 /CLONE=HEMBA1003989 /UG=Hs.274266 Homo sapiens cDNA FLJ10175 fis, clone HEMBA	AU145135	1.97
glycogen synthase 2 (liver)	S70004	1.97
hypothetical protein FLJ23142	NM_024583	1.97
potassium voltage-gated channel, Shal-related subfamily, member 2	NM_012281	1.97
hypothetical protein FLJ21777	AK025430	1.97
KIAA1009 protein	AK023613	1.97
chromosome 6 open reading frame 10	NM_006781	1.97
spectrin, beta, non-erythrocytic 1	R85779	1.97
Consensus includes gb:BC004145.1 /DEF=Homo sapiens, clone IMAGE:3509783, mRNA, partial cds. /FEA=mRNA /PROD=Unknown (protein for IMAGE:3509783) /DB_XREF=gi:13278737 /L	BC004145	1.97
Consensus includes gb:AL049390.1 /DEF=Homo sapiens mRNA; cDNA DKFZp58601318 (from clone DKFZp58601318). /FEA=mRNA /DB_XREF=gi:4500184 /UG=Hs.22689 Homo sapiens r	AL049390	1.97
Consensus includes gb:AW025579 /FEA=EST /DB_XREF=gi:5879109 /DB_XREF=est:wu75b04.x1 /CLONE=IMAGE:2525839 /UG=Hs.100261 Homo sapiens mRNA; cDNA DKFZp564B222 (fro	AW025579	1.97
polymerase (RNA) III (DNA directed) (32kD)	NM_006467	1.97
5-hydroxytryptamine (serotonin) receptor 4	Y08756	1.97
ubiquitin-specific processing protease	NM_020903	1.97
KIAA0553 protein	AI673812	1.97
Consensus includes gb:AU147200 /FEA=EST /DB_XREF=gi:11008721 /DB_XREF=est:AU147200 /CLONE=HEMBB1002699 /UG=Hs.296698 Homo sapiens cDNA FLJ12105 fis, clone HEMBB	AU147200	1.97
KIAA1013 protein	AU145019	1.97
KIAA0878 protein	N21138	1.97
hypothetical protein PRO1942	NM_018610	1.97
hypothetical protein FLJ10159	NM_018013	1.96
hypothetical protein PRO2714	NM_018534	1.96
heterogeneous nuclear ribonucleoprotein U (scaffold attachment factor A)	D13413	1.96
beta-amyloid binding protein precursor	BF968960	1.96
KIAA0256 gene product	N52532	1.96
hypothetical protein FLJ23188	NM_024628	1.96
RelA-associated inhibitor	NM_006663	1.96
forkhead box G1A	NM_004471	1.96

3-phosphoinositide dependent protein kinase-1	AK026947	1.96
Consensus includes gb:AL080106.1 /DEF=Homo sapiens mRNA; cDNA DKFZp5660053 (from clone DKFZp5660053). /FEA=mRNA /DB_XREF=gi:5262531 /UG=Hs.306323 Homo sapiens mi	AL080106	1.96
gb:U43279.1 /DEF=Human nucleoporin nup 36 mRNA, complete cds. /FEA=mRNA /PROD=nup 36 /DB_XREF=gi:1565299 /FL=gb:U43279.1	U43279	1.96
endothelial differentiation, lysophosphatidic acid G-protein-coupled receptor, 2	AW269335	1.96
ATPase, Class V, type 10C	N35112	1.96
monoamine oxidase A	NM_000240	1.96
doublesex and mab-3 related transcription factor 1	NM_021951	1.96
peroxisomal biogenesis factor 11A	AL360141	1.96
chemokine (C-X3-C) receptor 1	U20350	1.96
inactivation escape 2	AW003516	1.96
collagen, type VIII, alpha 1	BE877796	1.96
phosphoribosyl pyrophosphate synthetase 2	NM_002765	1.96
Consensus includes gb:AW970584 /FEA=EST /DB_XREF=gi:8160429 /DB_XREF=est:EST382665 /UG=Hs.291033 ESTs	AW970584	1.96
serine/threonine kinase 11 (Peutz-Jeghers syndrome)	NM_000455	1.96
Consensus includes gb:AL096727.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434B104 (from clone DKFZp434B104). /FEA=mRNA /DB_XREF=gi:5419860 /UG=Hs.168249 Homo sapiens mI	AL096727	1.96
EphB4	BC004264	1.96
Consensus includes gb:AB007970.1 /DEF=Homo sapiens mRNA, chromosome 1 specific transcript KIAA0501. /FEA=mRNA /DB_XREF=gi:3413945 /UG=Hs.223020 Homo sapiens mRNA, chi	AB007970	1.95
glyceraldehyde-3-phosphate dehydrogenase	AL035604	1.95
DKFZP564A122 protein	NM_015626	1.95
Src-like-adaptor	U44403	1.95
transmembrane gamma-carboxyglutamic acid protein 4	NM_024081	1.95
NCK-associated protein 1	AK001291	1.95
cytochrome P450, subfamily IIC (mephenytoin 4-hydroxylase), polypeptide 19	X65962	1.95
Consensus includes gb:AK025094.1 /DEF=Homo sapiens cDNA: FLJ21441 fis, clone COL04422. /FEA=mRNA /DB_XREF=gi:10437541 /UG=Hs.269926 Homo sapiens cDNA: FLJ21441 fis, cl	AK025094	1.95
Consensus includes gb:AL117630.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434D115 (from clone DKFZp434D115). /FEA=mRNA /DB_XREF=gi:5912213 /UG=Hs.252723 ribosomal protei	AL117630	1.95
hematopoietic PBX-interacting protein	NM_020524	1.95
hypothetical protein MGC12262	BC005236	1.95
KIAA1009 protein	NM_014895	1.95
oxidative stress induced like	U46752	1.95
BRCA1 associated protein	AL042733	1.95
RAB31, member RAS oncogene family	BE789881	1.95
gamma-aminobutyric acid (GABA) receptor, rho 1	NM_002042	1.95
hypothetical protein MGC4655	BC004908	1.95
integrin-binding sialoprotein (bone sialoprotein, bone sialoprotein II)	NM_004967	1.95
hypothetical protein FLJ10512	NM_018121	1.95
hypothetical protein FLJ20666	NM_018333	1.95
CD24 antigen (small cell lung carcinoma cluster 4 antigen)	M58664	1.95
KIAA1037 protein	AB028960	1.95
decorin	AF138303	1.95
ATP-binding cassette, sub-family B (MDR/TAP), member 11	AF136523	1.95
NK3 transcription factor homolog A (Drosophila)	AF249669	1.95
Consensus includes gb:AI679024 /FEA=EST /DB_XREF=gi:4889206 /DB_XREF=est:tu71a10.x1 /CLONE=IMAGE:2256474 /UG=Hs.201420 ESTs, Moderately similar to ALU7_HUMAN ALU SU	AI679024	1.95
hairy/enhancer-of-split related with YRPW motif 1	NM_012258	1.95
gb:AF165312.1 /DEF=Homo sapiens pre-T-cell receptor alpha chain (PTA) mRNA, alternatively spliced, complete cds. /FEA=CDS /GEN=PTA /PROD=pre-T-cell receptor alpha chain /DB_XRE	AF165312	1.94
a disintegrin-like and metalloprotease (reprolysin type) with thrombospondin type 1 motif, 1	AK023795	1.94
synaptonemal complex protein 1	X95654	1.94
KIAA0874 protein	X80821	1.94
chromosome 21 open reading frame 2	NM_004928	1.94
nudix (nucleoside diphosphate linked moiety X)-type motif 7	AK026469	1.94
hypothetical protein FLJ10648	AL049394	1.94
transcription elongation factor B (SIII), polypeptide 2 (18kD, elongin B)	AC004493	1.94
neurofilament 3 (150kD medium)	NM_005382	1.94
estrogen receptor 2 (ER beta)	AF060555	1.94
membrane-spanning 4-domains, subfamily A, member 2 (Fc fragment of IgE, high affinity I, receptor for; beta polypeptide)	NM_000139	1.94
olfactory receptor, family 1, subfamily G, member 1	NM_003555	1.94
Consensus includes gb:AK021571.1 /DEF=Homo sapiens cDNA FLJ11509 fis, clone HEMBA1002166. /FEA=mRNA /DB_XREF=gi:10432777 /UG=Hs.289093 Homo sapiens cDNA FLJ11509 fi	AK021571	1.94
thyroid hormone receptor interactor 10	NM_004240	1.94
Consensus includes gb:AK022280.1 /DEF=Homo sapiens cDNA FLJ12218 fis, clone MAMMA1001075, moderately similar to Homo sapiens CGI-72 protein mRNA. /FEA=mRNA /DB_XREF=gi:	AK022280	1.94
translation initiation factor IF2	BE138647	1.94
DKFZP564J102 protein	AL080065	1.94
nudix (nucleoside diphosphate linked moiety X)-type motif 7	AK026469	1.94
keratin 5 (epidermolysis bullosa simplex, Dowling-Meara/Kobner/Weber-Cockayne types)	NM_000424	1.94
glutamate receptor, ionotropic, N-methyl D-aspartate 2B	U11287	1.94

carbonic anhydrase XI	AB018195	1.94
prion protein (p27-30) (Creutzfeld-Jakob disease, Gerstmann-Strausler-Scheinker syndrome, fatal familial insomnia)	NM_000311	1.94
synapsin III	NM_003490	1.93
sparc/osteonectin, cwcv and kazal-like domains proteoglycan (testican)	AF231124	1.93
mucin and cadherin-like	NM_017717	1.93
zinc finger protein 42 (myeloid-specific retinoic acid- responsive)	M58297	1.93
aldehyde oxidase 1	AB046692	1.93
cholesterol 25-hydroxylase	NM_003956	1.93
angiotensin receptor 1	NM_004835	1.93
phosducin	M33478	1.93
retinoid X receptor, alpha	NM_002957	1.93
guanine nucleotide binding protein (G protein), alpha 11 (Gq class)	AI040021	1.93
integrin, alpha M (complement component receptor 3, alpha; also known as CD11b (p170), macrophage antigen alpha polypeptide)	NM_000632	1.93
heat shock 70kD protein 1A	NM_005345	1.93
alkaline phosphatase, intestinal	NM_001631	1.93
synaptojanin 1	NM_003895	1.93
AND-1 protein	AK001538	1.93
natriuretic peptide receptor C/guanylate cyclase C (atrionatriuretic peptide receptor C)	NM_000908	1.93
human epididymis-specific 3 alpha	NM_006683	1.93
oligodendrocyte lineage transcription factor 2	AA757419	1.93
chromosome 2 open reading frame 3	BC000853	1.93
matrix metalloproteinase 7 (matrilysin, uterine)	NM_002423	1.93
helicase KIAA0054	NM_014877	1.93
MAD, mothers against decapentaplegic homolog 5 (Drosophila)	AF010601	1.93
skin-specific protein	NM_014357	1.93
hypothetical protein PRO1496	NM_018603	1.92
transmembrane 6 superfamily member 2	AK024515	1.92
protocadherin alpha 9	NM_014005	1.92
KIAA0870 protein	AB020677	1.92
synaptotagmin XIII	AW089736	1.92
hypothetical protein MGC11138	BC004504	1.92
WW domain binding protein 4 (formin binding protein 21)	NM_007187	1.92
Consensus includes gb:AK021983.1 /DEF=Homo sapiens cDNA FLJ11921 fis, clone HEMBB1000318. /FEA=mRNA /DB_XREF=gi:10433292 /UG=Hs.306616 Homo sapiens cDNA FLJ11921 f	AK021983	1.92
Consensus includes gb:AL121981 /DEF=Human DNA sequence from clone RP5-1061C18 on chromosome 1p36.22-36.33. Contains ESTs, STSs, GSSs and a CpG island. Contains the 5 part of	AL121981	1.92
dopamine receptor D3	L20469	1.92
CD6 antigen	NM_006725	1.92
X03453 Bacteriophage P1 cre recombinase protein (-5 and -3 represent transcript regions 5 prime and 3 prime respectively)	X03453	1.92
NY-REN-58 antigen	NM_016122	1.92
chitinase 3-like 1 (cartilage glycoprotein-39)	M80927	1.92
glycine receptor, alpha 3	AW298713	1.92
Consensus includes gb:AK021495.1 /DEF=Homo sapiens cDNA FLJ11433 fis, clone HEMBA1001121. /FEA=mRNA /DB_XREF=gi:10432690 /UG=Hs.287415 Homo sapiens cDNA FLJ11433 fi	AK021495	1.92
Consensus includes gb:AK024602.1 /DEF=Homo sapiens cDNA: FLJ20949 fis, clone ADSE01902. /FEA=mRNA /DB_XREF=gi:10436915 /UG=Hs.220255 Homo sapiens cDNA: FLJ20949 fis,	AK024602	1.92
zinc finger protein 93 homolog (mouse)	BC002663	1.92
CGI-43 protein	NM_015622	1.92
double homeobox, 4	D38024	1.92
radixin	AJ057093	1.92
LIM domain protein	AF153882	1.92
RAN binding protein 1	AJ221318	1.92
small nuclear ribonucleoprotein polypeptide E	W87901	1.92
translocase of inner mitochondrial membrane 9 homolog (yeast)	NM_012460	1.92
DnaJ (Hsp40) homolog, subfamily C, member 4	AW024467	1.92
neurexophilin 3	AI937333	1.92
G-2 and S-phase expressed 1	BC006325	1.92
glutamate receptor, ionotropic, AMPA 1	AL567302	1.92
fibronectin 1	AJ276395	1.91
arachidonate 15-lipoxygenase	NM_001140	1.91
cytochrome P450, subfamily IIIA (niphedipine oxidase), polypeptide 4	NM_017460	1.91
hypothetical protein FLJ10430	NM_018092	1.91
Consensus includes gb:X61070.1 /DEF=Human mRNA for T cell receptor, clone IGRA15. /FEA=mRNA /PROD=T cell receptor /DB_XREF=gi:33508 /UG=Hs.121492 Human mRNA for T cell re	X61070	1.91
tripartite motif-containing 10	AF220122	1.91
variable charge, Y chromosome	NM_004679	1.91
cytochrome P450, subfamily XIB (steroid 11-beta-hydroxylase), polypeptide 2	X54741	1.91
protein phosphatase 2, regulatory subunit B (B56), alpha isoform	AI803525	1.91

MAD, mothers against decapentaplegic homolog 5 (Drosophila)	NM_005903	1.91
pregnancy specific beta-1-glycoprotein 1	M33663	1.91
BLu protein	NM_015896	1.91
histatin 1	NM_002159	1.91
DKFZP5640092 protein	AK022215	1.91
leucine zipper, down-regulated in cancer 1	NM_012317	1.91
KIAA0446 gene product	AB007915	1.91
calcium channel, voltage-dependent, alpha 1H subunit	NM_021098	1.91
killer cell lectin-like receptor subfamily D, member 1	NM_007334	1.91
Fc fragment of IgE, high affinity I, receptor for; alpha polypeptide	BC005912	1.91
hypothetical protein FLJ10650	NM_018168	1.91
Consensus includes gb:AA491625 /FEA=EST /DB_XREF=gi:2221187 /DB_XREF=est:ne80d01.s1 /CLONE=IMAGE:910561 /UG=Hs.293940 ESTs	AA491625	1.90
hypothetical protein FLJ20605	AL136931	1.90
Consensus includes gb:AK000156.1 /DEF=Homo sapiens cDNA FLJ20149 fis, clone COL08213. /FEA=mRNA /DB_XREF=gi:7020059 /UG=Hs.272193 Homo sapiens cDNA FLJ20149 fis, clon	AK000156	1.90
fatty acid binding protein 3, muscle and heart (mammary-derived growth inhibitor)	NM_004102	1.90
nephrosis 1, congenital, Finnish type (nephrin)	NM_004646	1.90
heat shock transcription factor 2 binding protein	BF056462	1.90
interleukin 21	NM_021803	1.90
VASA protein	AY004154	1.90
Consensus includes gb:AK025065.1 /DEF=Homo sapiens cDNA: FLJ21412 fis, clone COL03987. /FEA=mRNA /DB_XREF=gi:10437504 /UG=Hs.288066 Homo sapiens cDNA: FLJ21412 fis, cl	AK025065	1.90
tumor protein 63 kDa with strong homology to p53	AF116771	1.90
C9orf10 protein	AF214738	1.90
vacuolar protein sorting 45A (yeast)	NM_007258	1.90
myelin-associated oligodendrocyte basic protein	D28114	1.90
hypothetical protein	AA835004	1.90
thrombospondin 2	NM_003247	1.90
tyrosine aminotransferase	NM_000353	1.90
origin recognition complex, subunit 5-like (yeast)	AF081459	1.90
metaphase chromosome protein 1	NM_017513	1.90
5' nucleotidase, mitochondrial	NM_020201	1.90
angiopoietin 2	NM_001147	1.90
cadherin-like 22	AF035300	1.90
M11167 Human 28S rRNA sequence, length 5025 bases, middle target bases 1666-3330	M11167	1.90
gb:L34164.1 /DEF=Human Ig rearranged mu-chain gene VH3-D2110-JH2, complete cds., polyreactive. /FEA=mRNA /DB_XREF=gi:576594 /FL=gb:L34164.1	L34164	1.90
nuclear factor I/B	AJ700518	1.90
immunoglobulin lambda joining 3	AF234254	1.90
RAB3A, member RAS oncogene family	AA988241	1.90
M24537B subtilis pheB, pheA genes corresponding to nucleotides 2017-3334 of M24537 (-5, -M, -3 represent transcript regions 5 prime, Middle, and 3 prime respectively)	M24537B	1.90
fibrillin 1 (Marfan syndrome)	AJ264196	1.90
a disintegrin and metalloproteinase domain 22	AF155382	1.90
gb:AF274945.1 /DEF=Homo sapiens PNAS-20 mRNA, complete cds. /FEA=mRNA /PROD=PNAS-20 /DB_XREF=gi:12751054 /FL=gb:AF274945.1	AF274945	1.90
KIAA1536 protein	BC003177	1.90
hypothetical protein MGC5499	NM_024055	1.90
G protein-coupled receptor 3	AW024085	1.90
hypothetical protein FLJ21625	NM_025039	1.90
UDP-Gal:betaGlcNAc beta 1,3-galactosyltransferase, polypeptide 2	Y15014	1.90
neuromedin B receptor	NM_002511	1.90
ankyrin repeat and SOCS box-containing 4	AC003079	1.90
formyl peptide receptor 1	NM_002029	1.89
solute carrier family 6 (neurotransmitter transporter, taurine), member 6	BC006252	1.89
UDP-N-acetylglucosamine:a-1,3-D-mannoside beta-1,4-N-acetylglucosaminyltransferase IV-homolog	NM_013244	1.89
Consensus includes gb:AF131813.1 /DEF=Homo sapiens clone 24970 mRNA sequence. /FEA=mRNA /DB_XREF=gi:4406647 /UG=Hs.21258 Homo sapiens cDNA FLJ13094 fis, clone NT2RP	AF131813	1.89
lymphoid nuclear protein related to AF4	NM_002285	1.89
arylsulfatase D	NM_009589	1.89
hypothetical protein FLJ11348	NM_018395	1.89
Consensus includes gb:AK000145.1 /DEF=Homo sapiens cDNA FLJ20138 fis, clone COL07172. /FEA=mRNA /DB_XREF=gi:7020041 /UG=Hs.306384 Homo sapiens cDNA FLJ20138 fis, clon	AK000145	1.89
similar to rat nuclear ubiquitous casein kinase 2	AW515443	1.89
M10098 Human 18S rRNA gene, complete (_5, _M, _3 represent transcript regions 5 prime, Middle, and 3 prime respectively)	M10098	1.89
hypothetical protein dJ462O23.2	BC001265	1.89
microtubule-associated protein 1A	AA633627	1.89
hypothetical gene C6O18	AL049786	1.89
hypothetical protein FLJ13885	NM_025016	1.89
annexin A4	NM_001153	1.89

Consensus includes gb:BE467941 /FEA=EST /DB_XREF=gi:9513716 /DB_XREF=est:hz76b04.x1 /CLONE=IMAGE:3213871 /UG=Hs.250175 homolog of yeast long chain polyunsaturated fat	BE467941	1.89
solute carrier family 1 (glutamate transporter), member 7	NM_006671	1.89
tumor protein p53 (Li-Fraumeni syndrome)	K03199	1.89
nyctalopin	AF254868	1.89
Consensus includes gb:M18917 /DEF=Human glucocerebrosidase gene 10-2, 5 end /FEA=CDS_1 /DB_XREF=gi:183025 /UG=Hs.287396 Human glucocerebrosidase gene 10-2, 5 end	M18917	1.89
KIAA0451 gene product	AK027000	1.89
insulin-like growth factor binding protein 5	M65062	1.89
Sp3 transcription factor	AU145005	1.89
parvin, alpha	NM_018222	1.89
caspase 8, apoptosis-related cysteine protease	BF439983	1.89
zinc finger protein, subfamily 1A, 4 (Eos)	NM_022465	1.89
KIAA0138 gene product	NM_014649	1.89
p21 (CDKN1A)-activated kinase 2	NM_002577	1.89
SET translocation (myeloid leukemia-associated)	W26593	1.88
KIAA1659 protein	AB051446	1.88
hypothetical protein MGC3771	NM_030970	1.88
hypothetical protein FLJ14326	AK024388	1.88
growth differentiation factor 10	NM_004962	1.88
ALEX3 protein	NM_016607	1.88
v-akt murine thymoma viral oncogene homolog 2	M77198	1.88
Consensus includes gb:AI825798 /FEA=EST /DB_XREF=gi:5446469 /DB_XREF=est:td18e08.x1 /CLONE=IMAGE:2076038 /UG=Hs.211388 Homo sapiens BAC clone CTB-60N22 from 7q21	AI825798	1.88
Consensus includes gb:AK024553.1 /DEF=Homo sapiens cDNA: FLJ20900 fis, clone ADKA03593. /FEA=mRNA /DB_XREF=gi:10436861 /UG=Hs.306687 Homo sapiens cDNA: FLJ20900 fis,	AK024553	1.88
glutathione peroxidase 3 (plasma)	AW149846	1.88
lamin A/C	NM_005572	1.88
guanine nucleotide binding protein (G protein), alpha 15 (Gq class)	NM_002068	1.88
Consensus includes gb:AW974823 /FEA=EST /DB_XREF=gi:8166026 /DB_XREF=est:EST386928 /UG=Hs.105411 ESTs	AW974823	1.88
gb:BC001407.1 /DEF=Homo sapiens, Similar to cytochrome c-like antigen, clone MGC:2960, mRNA, complete cds. /FEA=mRNA /PROD=Similar to cytochrome c-like antigen /DB_XREF=gi:12	BC001407	1.88
hydroxysteroid (17-beta) dehydrogenase 2	NM_002153	1.88
non-metastatic cells 4, protein expressed in	AL523860	1.88
hypothetical protein FLJ12876	NM_022754	1.88
CD5 antigen-like (scavenger receptor cysteine rich family)	NM_005894	1.88
cold autoinflammatory syndrome 1	AK027194	1.88
hypothetical protein FLJ14033 similar to hypoxia inducible factor 3, alpha subunit	AK021881	1.88
hypothetical protein FLJ22490	NM_024790	1.88
hypothetical protein FLJ20081	NM_017658	1.88
aquaporin 1 (channel-forming integral protein, 28kD)	AL518391	1.88
RAR-related orphan receptor A	U04897	1.88
pleckstrin homology-like domain, family A, member 3	NM_012396	1.88
solute carrier family 16 (monocarboxylic acid transporters), member 7	NM_004731	1.88
S100 calcium binding protein A12 (calgranulin C)	NM_005621	1.88
bullous pemphigoid antigen 1 (230/240kD)	AL096710	1.88
amyloid beta (A4) precursor-like protein 2	BC000373	1.88
POU domain, class 4, transcription factor 1	NM_006237	1.88
hypothetical protein FLJ22332	NM_024724	1.87
lecithin retinol acyltransferase (phosphatidylcholine--retinol O-acyltransferase)	NM_004744	1.87
KIAA0798 gene product	NM_014650	1.87
hypothetical protein PRO2266	NM_018519	1.87
hypothetical protein MGC12488	BC005372	1.87
similar to rat myomegalin	BC005949	1.87
alcohol dehydrogenase 1C (class I), gamma polypeptide	NM_000669	1.87
amyloid beta (A4) precursor protein (protease nexin-II, Alzheimer disease)	NM_000484	1.87
trinucleotide repeat containing 15	AB014542	1.87
hypothetical protein FLJ22324	NM_024120	1.87
zinc finger protein 291	AF119814	1.87
fibroblast growth factor 12	AL119322	1.87
6 protein-coupled receptor 44	NM_004778	1.87
Consensus includes gb:AL121890 /DEF=Human DNA sequence from clone RP5-1116H23 on chromosome 20 Contains a novel gene, a 40S ribosomal protein S21 pseudogene, 2 CpG islands,	AL121890	1.87
transmembrane protease, serine 5 (spinesin)	NM_030770	1.87
transferrin	AJ073407	1.87
methionine adenosyltransferase II, alpha	AW301861	1.87
like mouse brain protein E46	Z95331	1.87
protein kinase, cAMP-dependent, catalytic, gamma	NM_002732	1.87
erythropoietin receptor	M34986	1.87

spectrin, alpha, erythrocytic 1 (elliptocytosis 2)	NM_003126	1.87
nuclear factor of kappa light polypeptide gene enhancer in B-cells 2 (p49/p100)	U09609	1.87
Consensus includes gb:AK021609.1 /DEF=Homo sapiens cDNA FLJ11547 fis, clone HEMBA1002934. /FEA=mRNA /DB_XREF=gi:10432824 /UG=Hs.306605 Homo sapiens cDNA FLJ11547 fi	AK021609	1.87
lunatic fringe homolog (Drosophila)	U94354	1.87
GRO2 oncogene	M57731	1.87
Consensus includes gb:AA131826 /FEA=EST /DB_XREF=gi:1693379 /DB_XREF=est:zl37f06.s1 /CLONE=IMAGE:504131 /UG=Hs.107164 spectrin, beta, non-erythrocytic 1	AA131826	1.87
hypothetical protein FLJ12076	NM_025187	1.87
hypothetical protein FLJ13590	NM_024840	1.86
golgi associated, gamma adaptin ear containing, ARF binding protein 1	AW001443	1.86
similar to rab11-binding protein	NM_019045	1.86
gb:AF034374.1 /DEF=Homo sapiens molybdenum cofactor biosynthesis protein A and molybdenum cofactor biosynthesis protein C mRNA, complete cds. /FEA=mRNA /PROD=molybdenum	AF034374	1.86
Consensus includes gb:AK023843.1 /DEF=Homo sapiens cDNA FLJ13781 fis, clone PLACE4000465. /FEA=mRNA /DB_XREF=gi:10435902 /UG=Hs.252820 Homo sapiens cDNA FLJ13781 fi	AK023843	1.86
kynureninase (L-kynurenine hydrolase)	BC000879	1.86
peripheral benzodiazepine receptor-associated protein 1	NM_004758	1.86
KIAA0738 gene product	NM_014719	1.86
PRO0650 protein	NM_014137	1.86
hypothetical protein FLJ20783	W87466	1.86
Consensus includes gb:AI824078 /FEA=EST /DB_XREF=gi:5444749 /DB_XREF=est:wj35h06.x1 /CLONE=IMAGE:2404859 /UG=Hs.295446 ESTs, Moderately similar to 810024C cytochrome	AI824078	1.86
protocadherin beta 12	NM_018932	1.86
fibroblast growth factor receptor 2 (bacteria-expressed kinase, keratinocyte growth factor receptor, craniofacial dysostosis 1, Crouzon syndrome, Pfeiffer syndrome, Jackson-Weiss syndrome	NM_022969	1.86
signal transducer and activator of transcription 4	NM_003151	1.86
chromosome 5 open reading frame 7	NM_016604	1.86
Consensus includes gb:AK023838.1 /DEF=Homo sapiens cDNA FLJ13776 fis, clone PLACE4000387. /FEA=mRNA /DB_XREF=gi:10435896 /UG=Hs.287595 Homo sapiens cDNA FLJ13776 fi	AK023838	1.86
olfactory receptor, family 10, subfamily H, member 1	NM_013940	1.86
basement membrane-induced gene	AB035482	1.86
Smcx homolog, X chromosome (mouse)	NM_004187	1.86
hypothetical protein similar to rat vacuolar sorting protein 33a	NM_022916	1.86
hypothetical protein MGC3036	NM_023942	1.86
zinc finger protein 345	NM_003419	1.86
leucine-rich repeat-containing 5	NM_018103	1.86
Consensus includes gb:AL137312.1 /DEF=Homo sapiens mRNA; cDNA DKFZp761K23121 (from clone DKFZp761K23121). /FEA=mRNA /DB_XREF=gi:6807797 /UG=Hs.293681 Homo sapie	AL137312	1.86
Consensus includes gb:AK021967.1 /DEF=Homo sapiens cDNA FLJ11905 fis, clone HEMBB1000050. /FEA=mRNA /DB_XREF=gi:10433276 /UG=Hs.273759 Homo sapiens cDNA FLJ11905 f	AK021967	1.86
empty spiracles homolog 1 (Drosophila)	AW196403	1.86
Consensus includes gb:AL033403 /DEF=Human DNA sequence from clone 88D7 on chromosome Xq25-26.3 Contains F9 (coagulation factor IX (plasma thromboplastic component, Christmas	AL033403	1.86
solute carrier family 16 (monocarboxylic acid transporters), member 4	NM_004696	1.86
endothelin receptor type B	NM_000115	1.85
H2B histone family, member E	NM_003521	1.85
sodium channel, voltage-gated, type III, alpha polypeptide	AF225986	1.85
leucine zipper, putative tumor suppressor 1	AF123654	1.85
zinc finger protein 58ZF3	NM_020394	1.85
hypothetical protein FLJ22684	NM_025048	1.85
chemokine (C-C motif) receptor 1	NM_001295	1.85
laminin, gamma 2 (nicein (100kD), kalinin (105kD), BM600 (100kD), Herlitz junctional epidermolysis bullosa))	NM_005562	1.85
Consensus includes gb:AK024606.1 /DEF=Homo sapiens cDNA: FLJ20953 fis, clone ADSE01979. /FEA=mRNA /DB_XREF=gi:10436920 /UG=Hs.306697 Homo sapiens cDNA: FLJ20953 fis,	AK024606	1.85
forkhead box L1	AF315075	1.85
hypothetical protein FLJ11783	NM_024891	1.85
U2 small nuclear ribonucleoprotein auxiliary factor (65kD)	AI810156	1.85
gb:AF202635.1 /DEF=Homo sapiens PP1200 mRNA, complete cds. /FEA=mRNA /PROD=PP1200 /DB_XREF=gi:10732645 /UG=Hs.302135 Homo sapiens PP1200 mRNA, complete cds /FL=	AF202635	1.85
Consensus includes gb:AL080190.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434A202 (from clone DKFZp434A202). /FEA=mRNA /DB_XREF=gi:5262670 /UG=Hs.189242 Homo sapiens mI	AL080190	1.85
debranching enzyme homolog 1 (S. cerevisiae)	NM_016216	1.85
EST-YD1 protein	NM_021208	1.85
mucin 5, subtype B, tracheobronchial	AI697108	1.85
carbonic anhydrase III, muscle specific	NM_005181	1.85
ceruloplasmin (ferroxidase)	NM_000096	1.85
Cluster Incl. AI937160:wp73h02.x1 Homo sapiens cDNA, 3' end /clone=IMAGE-2467443 /clone_end=3' /gb=AI937160 /gi=5676030 /ug=Hs.7967 /len=552	AI937160	1.85
colony stimulating factor 1 (macrophage)	M76453	1.85
neural retina leucine zipper	M81840	1.85
Consensus includes gb:AL049252.1 /DEF=Homo sapiens mRNA; cDNA DKFZp564D193 (from clone DKFZp564D193). /FEA=mRNA /DB_XREF=gi:4499993 /UG=Hs.302048 Homo sapiens mI	AL049252	1.85
B-cell CLL/lymphoma 2	M13994	1.85
PRO0456 protein	NM_014127	1.85
platelet-activating factor receptor	D10202	1.85
mitogen-activated protein kinase kinase kinase kinase 5	Z25426	1.85
lymphocyte antigen 9	AF244129	1.85

LIM domain protein	BE043700	1.85
RFamide-related peptide precursor	NM_022150	1.84
Consensus includes gb:AK021457.1 /DEF=Homo sapiens cDNA FLJ11395 fis, clone HEMBA1000594. /FEA=mRNA /DB_XREF=gi:10432649 /UG=Hs.306595 Homo sapiens cDNA FLJ11395 f	AK021457	1.84
H3 histone family, member K	NM_003536	1.84
CD6 antigen	U66145	1.84
placental growth factor, vascular endothelial growth factor-related protein	BC001422	1.84
fer-1-like 3, myoferlin (C. elegans)	NM_013451	1.84
chymase 1, mast cell	NM_001836	1.84
carbohydrate kinase-like	NM_013276	1.84
hypothetical protein	NM_019079	1.84
acetyl LDL receptor; SREC=scavenger receptor expressed by endothelial cells	NM_003693	1.84
frizzled homolog 1 (Drosophila)	AF072872	1.84
homeo box D3	Y09980	1.84
aldolase A, fructose-bisphosphate pseudogene 2	M21191	1.84
CD24 antigen (small cell lung carcinoma cluster 4 antigen)	X69397	1.84
6 protein-coupled receptor kinase 2-like (Drosophila)	U33054	1.84
nuclear transcription factor Y, alpha	AL031778	1.84
6 protein-coupled receptor 57	NM_014627	1.84
PC3-96 protein	NM_022488	1.84
interleukin 1, zeta	NM_014439	1.84
Consensus includes gb:AL109667.1 /DEF=Homo sapiens mRNA full length insert cDNA clone EUROIMAGE 381867. /FEA=mRNA /DB_XREF=gi:5689815 /UG=Hs.124154 Homo sapiens mRN	AL109667	1.84
major histocompatibility complex, class II, DQ beta 1	M24364	1.84
protocadherin 9	NM_020403	1.84
cadherin 17, LI cadherin (liver-intestine)	U07969	1.84
keratin 14 (epidermolysis bullosa simplex, Dowling-Meara, Koebner)	BC002690	1.84
KIAA0256 gene product	D87445	1.84
bromodomain adjacent to zinc finger domain, 1A	AA102574	1.84
uroplakin 1B	NM_006952	1.84
Consensus includes gb:AW451230 /FEA=EST /DB_XREF=gi:6992006 /DB_XREF=est:UI-H-BI3-alh-a-01-0-UI.s1 /CLONE=IMAGE:2736649 /UG=Hs.257332 ESTs, Highly similar to KIAA0311 I	AW451230	1.84
Consensus includes gb:AL109716.2 /DEF=Homo sapiens mRNA full length insert cDNA clone EUROIMAGE 208948. /FEA=mRNA /DB_XREF=gi:9187597 /UG=Hs.20034 Homo sapiens mRNA	AL109716	1.84
male germ cell-associated kinase	NM_005906	1.84
glycoprotein VI (platelet)	AB043821	1.84
KIAA0865 protein	AJ522028	1.84
Consensus includes gb:AK022408.1 /DEF=Homo sapiens cDNA FLJ12346 fis, clone MAMMA1002297, highly similar to Homo sapiens mRNA for Rab6 GTPase activating protein. /FEA=mRNA	AK022408	1.84
homeo box D13	NM_000523	1.83
KIAA0974 protein	AB023191	1.83
serine (or cysteine) proteinase inhibitor, clade A (alpha-1 antiproteinase, antitrypsin), member 10	NM_016186	1.83
protein phosphatase 1, regulatory (inhibitor) subunit 1A	NM_006741	1.83
KIAA0643 protein	AA207013	1.83
A kinase (PRKA) anchor protein 6	NM_004274	1.83
hematopoietically expressed homeobox	NM_001529	1.83
chitinase 3-like 1 (cartilage glycoprotein-39)	AJ251847	1.83
Consensus includes gb:AU147295 /FEA=EST /DB_XREF=gi:11008816 /DB_XREF=est:AU147295 /CLONE=MAMMA1000264 /UG=Hs.288575 Homo sapiens cDNA FLJ14090 fis, clone MAMM	AU147295	1.83
hypothetical protein MGC14376	AF070569	1.83
calcium/calmodulin-dependent protein kinase (CaM kinase) II alpha	NM_015981	1.83
Consensus includes gb:AK025325.1 /DEF=Homo sapiens cDNA: FLJ21672 fis, clone COL09025. /FEA=mRNA /DB_XREF=gi:10437818 /UG=Hs.306801 Homo sapiens cDNA: FLJ21672 fis, cl	AK025325	1.83
RAB5B, member RAS oncogene family	AF267863	1.83
kinesin family member 13A	NM_022113	1.83
cytochrome P450, subfamily IIC (mephenytoin 4-hydroxylase), polypeptide 9	AV646536	1.83
Consensus includes gb:AI570199 /FEA=EST /DB_XREF=gi:4533573 /DB_XREF=est:to76b07.x1 /CLONE=IMAGE:2184181 /UG=Hs.182183 Homo sapiens mRNA for caldesmon, 3 UTR	AI570199	1.83
cell division cycle 25B	AL109804	1.83
SRY (sex determining region Y)-box 11	AW157202	1.83
KIAA0599 protein	BC004298	1.83
integrin, alpha 2b (platelet glycoprotein IIb of IIb/IIIa complex, antigen CD41B)	NM_000419	1.83
hypothetical protein FLJ10044	NM_017980	1.83
hypothetical protein FLJ13605	NM_024577	1.83
Consensus includes gb:BF222916 /FEA=EST /DB_XREF=gi:11130093 /DB_XREF=est:7q25a04.x1 /CLONE=IMAGE:3699342 /UG=Hs.331207 ESTs	BF222916	1.83
Consensus includes gb:AL121582 /DEF=Human DNA sequence from clone RP11-28009 on chromosome 20. Contains an RNF11 (ring finger protein 11) (SID1669) pseudogene, ESTs, STSs	AL121582	1.83
achaete-scute complex-like 1 (Drosophila)	BC001638	1.83
Consensus includes gb:AW973834 /FEA=EST /DB_XREF=gi:8165022 /DB_XREF=est:EST385936 /UG=Hs.105884 ESTs	AW973834	1.83
Consensus includes gb:AK000244.1 /DEF=Homo sapiens cDNA FLJ20237 fis, clone COLF5879, highly similar to AB023230 Homo sapiens mRNA for KIAA1013 protein. /FEA=mRNA /DB_XRE	AK000244	1.83
bullous pemphigoid antigen 1 (230/240kD)	NM_001723	1.83
bromodomain, testis-specific	NM_001726	1.83

proteasome (prosome, macropain) subunit, beta type, 2	BC000268	1.83
hypothetical protein FLJ11871	NM_025117	1.83
G-substrate	NM_006658	1.83
Consensus includes gb:AK023798.1 /DEF=Homo sapiens cDNA FLJ13736 fis, clone PLACE3000156. /FEA=mRNA /DB_XREF=gi:10435842 /UG=Hs.31532 Homo sapiens mRNA; cDNA DKFZp	AK023798	1.83
hypothetical protein FLJ11637	NM_024967	1.83
DNA-dependent protein kinase catalytic subunit-interacting protein 2	AI478592	1.83
endomucin-2	NM_016242	1.83
dystrobrevin, alpha	U26744	1.83
5-hydroxytryptamine (serotonin) receptor 7 (adenylate cyclase-coupled)	NM_019859	1.82
contactin 2 (axonal)	NM_005076	1.82
beta-site APP-cleaving enzyme	NM_012104	1.82
nuclear receptor subfamily 4, group A, member 2	NM_006186	1.82
complement factor H-related 4	NM_006684	1.82
glutamate receptor, ionotropic, AMPA 1	M64752	1.82
Consensus includes gb:BF674749 /FEA=EST /DB_XREF=gi:11948644 /DB_XREF=est:602136893F1 /CLONE=IMAGE:4273270 /UG=Hs.258800 ESTs	BF674749	1.82
Consensus includes gb:AK023446.1 /DEF=Homo sapiens cDNA FLJ13384 fis, clone PLACE1001062, highly similar to Homo sapiens mRNA for lysine-ketoglutarate reductasesaccharopine deh	AK023446	1.82
GIOT-2 for gonadotropin inducible transcription repressor-2	NM_016264	1.82
olfactory receptor, family 7, subfamily A, member 17	AC005255	1.82
visual system homeobox 1 homolog, CHX10-like (zebrafish)	NM_014588	1.82
adenylate cyclase 2 (brain)	AB028983	1.82
KIAA1031 protein	NM_025013	1.82
Consensus includes gb:AI693193 /FEA=EST /DB_XREF=gi:4970533 /DB_XREF=est:wd68f02.x1 /CLONE=IMAGE:2336763 /UG=Hs.247551 metaxin 1	AI693193	1.82
solute carrier family 11 (proton-coupled divalent metal ion transporters), member 1	D50402	1.82
immunoglobulin kappa constant	BG482805	1.82
chorionic gonadotropin, beta polypeptide	NM_000737	1.82
muscle, skeletal, receptor tyrosine kinase	NM_005592	1.82
gb:NM_022155.1 /DEF=Homo sapiens PP3227 protein (PP3227), mRNA. /FEA=mRNA /GEN=PP3227 /PROD=PP3227 protein /DB_XREF=gi:11545901 /UG=Hs.302027 PP3227 protein /F	NM_022155	1.82
hypothetical protein FLJ22297	NM_025135	1.82
nuclear receptor subfamily 0, group B, member 2	NM_021969	1.82
likely ortholog of mouse early B-cell factor 2	NM_022659	1.82
hypothetical protein FLJ11336	NM_018393	1.82
GRB2-related adaptor protein	NM_006613	1.82
insulin-like growth factor 2, antisense	NM_016412	1.82
synovial sarcoma translocation gene on chromosome 18-like 1	AB014593	1.82
hypothetical protein FLJ14117	NM_022777	1.82
CASP8 and FADD-like apoptosis regulator	AF009616	1.82
KIAA0680 gene product	AW295193	1.82
hypothetical protein FLJ10034	NM_017973	1.82
tumor necrosis factor (ligand) superfamily, member 15	NM_005118	1.82
hypothetical protein FLJ21269	NM_025107	1.82
egl nine homolog 3 (C. elegans)	NM_022073	1.82
solute carrier family 26, member 4	NM_000441	1.82
potassium inwardly-rectifying channel, subfamily J, member 9	NM_004983	1.82
sodium channel, voltage-gated, type IV, alpha polypeptide	NM_000334	1.82
hypothetical protein FLJ14084	NM_021637	1.82
Lin-7b protein; likely ortholog of mouse LIN-7B; mammalian LIN-7 protein 2	NM_022165	1.82
Glutamate receptor interacting protein	AI203397	1.82
KIAA0415 gene product	AI373854	1.82
myeloid/lymphoid or mixed-lineage leukemia (trithorax homolog, Drosophila); translocated to, 10	AF060938	1.82
Consensus includes gb:AW190208 /FEA=EST /DB_XREF=gi:6464688 /DB_XREF=est:zf60a12.x1 /CLONE=IMAGE:2679070 /UG=Hs.233789 ESTs	AW190208	1.82
chemokine ((-C motif) receptor 5	NM_000579	1.81
fibronectin leucine rich transmembrane protein 3	NM_013281	1.81
angiotensin receptor 2	NM_000686	1.81
PRO0038 protein	NM_014113	1.81
phosphoinositide-3-kinase, regulatory subunit, polypeptide 1 (p85 alpha)	AI934473	1.81
diacylglycerol kinase, beta (90kD)	AB018261	1.81
KIAA0672 gene product	AK023797	1.81
RBP1-like protein	AI051476	1.81
serine (or cysteine) proteinase inhibitor, clade B (ovalbumin), member 10	NM_005024	1.81
cyclic nucleotide gated channel beta 1	U18945	1.81
acetylcholinesterase (YT blood group)	M55040	1.81
cytokine receptor-like factor 3	NM_015986	1.81
Consensus includes gb:BF591638 /FEA=EST /DB_XREF=gi:11683962 /DB_XREF=est:7n53g03.x1 /CLONE=IMAGE:3568325 /UG=Hs.129663 ESTs	BF591638	1.81

Consensus includes gb:AK021576.1 /DEF=Homo sapiens cDNA FLJ11514 fis, clone HEMBA1002229. /FEA=mRNA /DB_XREF=gi:10432783 /UG=Hs.306602 Homo sapiens cDNA FLJ11514 f	AK021576	1.81
insulin-like growth factor binding protein 5	NM_000599	1.81
6 antigen 7	NM_021123	1.81
Consensus includes gb:AK023891.1 /DEF=Homo sapiens cDNA FLJ13829 fis, clone THYRO1000625. /FEA=mRNA /DB_XREF=gi:10435964 /UG=Hs.287600 Homo sapiens cDNA FLJ13829 fi	AK023891	1.81
F-box only protein 9	AL137520	1.81
KIAA1102 protein	AK026815	1.81
pancreatic lipase-related protein 1	NM_006229	1.81
cholinergic receptor, nicotinic, alpha polypeptide 1 (muscle)	BC006314	1.81
PR domain containing 2, with ZNF domain	U23736	1.81
hypothetical protein FLJ20556	NM_017878	1.81
phosphoinositol 3-phosphate-binding protein-2	BC000969	1.81
interleukin 13 receptor, alpha 2	NM_000640	1.81
H2A histone family, member O	BC001629	1.81
dihydropyrimidinase-like 4	AI857455	1.81
Consensus includes gb:AL050026.1 /DEF=Homo sapiens mRNA; cDNA DKFZp566C0224 (from clone DKFZp566C0224). /FEA=mRNA /DB_XREF=gi:4884265 /UG=Hs.306306 Homo sapiens	AL050026	1.81
complement component 9	K02766	1.81
Putative prostate cancer tumor suppressor	AU158251	1.81
hypothetical protein PRO1048	NM_018497	1.81
zinc finger protein 161	NM_007146	1.81
achaete-scute complex-like 1 (Drosophila)	BE797438	1.81
kynureninase (L-kynurenine hydrolase)	NM_003937	1.81
Snf2-related CBP activator protein	NM_006662	1.81
EphA5	L36644	1.81
myeloid/lymphoid or mixed-lineage leukemia (trithorax homolog, Drosophila); translocated to, 4	AB016898	1.80
Consensus includes gb:AL080130.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434E033 (from clone DKFZp434E033). /FEA=mRNA /DB_XREF=gi:5262569 /UG=Hs.15740 Homo sapiens mR	AL080130	1.80
KIAA1361 protein	AK024376	1.80
Consensus includes gb:AF052095.1 /DEF=Homo sapiens clone 23911 mRNA sequence. /FEA=mRNA /DB_XREF=gi:3360401 /UG=Hs.167344 Homo sapiens clone 23911 mRNA sequence	AF052095	1.80
zinc finger protein 79 (pT7)	X65232	1.80
leucine-rich repeat protein, neuronal 3	AI221950	1.80
calcium channel, voltage-dependent, gamma subunit 2	NM_006078	1.80
ATP-binding cassette, sub-family D (ALD), member 2	NM_005164	1.80
phosphodiesterase 10A	AB026816	1.80
myosin, heavy polypeptide 1, skeletal muscle, adult	NM_005963	1.80
karyopherin alpha 1 (importin alpha 5)	BC002374	1.80
cytlicin, basic protein of sperm head cytoskeleton 1	Z22780	1.80
Consensus includes gb:BG495771 /FEA=EST /DB_XREF=gi:13457287 /DB_XREF=est:602540396F1 /CLONE=IMAGE:4671599 /UG=Hs.289068 Homo sapiens cDNA FLJ11918 fis, clone HEI	BG495771	1.80
Consensus includes gb:AL137063 /DEF=Human DNA sequence from clone RP1-104L14 on chromosome 6 Contains part of the AKAP7 (A kinase (PRKA) anchor protein 7) gene, ESTs, STSs a	AL137063	1.80
BH-protocadherin (brain-heart)	AB006757	1.80
Consensus includes gb:AF009267.1 /DEF=Homo sapiens clone FBA1 Cri-du-chat region mRNA. /FEA=mRNA /DB_XREF=gi:2331069 /UG=Hs.102238 Homo sapiens clone FBA1 Cri-du-chat n	AF009267	1.80
hypothetical protein FLJ21135	NM_024866	1.80
paternally expressed 3	AF208967	1.80
mitogen-activated protein kinase kinase kinase 11	NM_002419	1.80
mab-21-like 2 (C. elegans)	AF262032	1.80
solute carrier family 4, sodium bicarbonate cotransporter, member 7	AF047033	1.80
bone morphogenetic protein receptor, type IA	AI678679	1.80
interleukin 8 receptor, beta	NM_001557	1.80
Consensus includes gb:AU147942 /FEA=EST /DB_XREF=gi:11009463 /DB_XREF=est:AU147942 /CLONE=MAMMA1002198 /UG=Hs.287517 Homo sapiens cDNA FLJ12333 fis, clone MAMM	AU147942	1.80
zinc finger protein 253	NM_021047	1.80
Consensus includes gb:AL117520.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434P2435 (from clone DKFZp434P2435). /FEA=mRNA /DB_XREF=gi:5912036 /UG=Hs.306348 Homo sapiens	AL117520	1.80
myosin IK	BE790157	1.80
insulin-like growth factor binding protein 2 (36kD)	NM_000597	1.80
one cut domain, family member 1	U96173	1.80
calpain 6	NM_014289	1.80
hormonally upregulated Neu-associated kinase	NM_014586	1.80
attractin	AL132773	1.80
KIAA1264 protein	AB040812	1.80
phosphate cytidylyltransferase 1, choline, beta isoform	NM_004845	1.80
KIAA0961 protein	NM_014898	1.80
Consensus includes gb:AK024615.1 /DEF=Homo sapiens cDNA: FLJ20962 fis, clone ADSH00804. /FEA=mRNA /DB_XREF=gi:10436931 /UG=Hs.306698 Homo sapiens cDNA: FLJ20962 fis,	AK024615	1.80
lysyl oxidase	L16895	1.80
Consensus includes gb:AU145357 /FEA=EST /DB_XREF=gi:11006878 /DB_XREF=est:AU145357 /CLONE=HEMBA1004611 /UG=Hs.296700 Homo sapiens cDNA FLJ12134 fis, clone MAMM/	AU145357	1.80
Consensus includes gb:AK026045.1 /DEF=Homo sapiens cDNA: FLJ22392 fis, clone HRC07868. /FEA=mRNA /DB_XREF=gi:10438761 /UG=Hs.275464 Homo sapiens cDNA: FLJ22392 fis, cl	AK026045	1.80
hypothetical protein FLJ10139	NM_018005	1.80

hypothetical protein PRO1728	NM_018505	1.80
Consensus includes gb:AL133396 /DEF=Human DNA sequence from clone RP5-1068H6 on chromosome 20q11.1-11.23. Contains a pseudogene similar to IDI1 (isopentenyl-diphosphate del	AL133396	1.80
microsomal triglyceride transfer protein (large polypeptide, 88kD)	AI623321	1.80
cathepsin D (lysosomal aspartyl protease)	AI560951	1.79
formyl peptide receptor-like 2	NM_002030	1.79
PCF11p homolog	AB020631	1.79
sarcoma antigen	NM_018666	1.79
ectodermal dysplasia 1, anhidrotic	AF061190	1.79
Consensus includes gb:AK022322.1 /DEF=Homo sapiens cDNA FLJ12260 fis, clone MAMMA1001551. /FEA=mRNA /DB_XREF=gi:10433693 /UG=Hs.306634 Homo sapiens cDNA FLJ12260 f	AK022322	1.79
POM (POM121 rat homolog) and ZP3 fusion	NM_012230	1.79
prothymosin, alpha (gene sequence 28)	AF348514	1.79
Consensus includes gb:AF009267.1 /DEF=Homo sapiens clone FBA1 Cri-du-chat region mRNA. /FEA=mRNA /DB_XREF=gi:2331069 /UG=Hs.102238 Homo sapiens clone FBA1 Cri-du-chat r	AF009267	1.79
GATA binding protein 6	D87811	1.79
Consensus includes gb:AU147851 /FEA=EST /DB_XREF=gi:11009372 /DB_XREF=est:AU147851 /CLONE=MAMMA1001878 /UG=Hs.183819 Homo sapiens cDNA FLJ12304 fis, clone MAMM	AU147851	1.79
Consensus includes gb:AL049342.1 /DEF=Homo sapiens mRNA; cDNA DKFZp566A193 (from clone DKFZp566A193). /FEA=mRNA /DB_XREF=gi:4500128 /UG=Hs.274506 Homo sapiens mf	AL049342	1.79
hypothetical protein FLJ12057	NM_024768	1.79
preproprolactin-releasing peptide	NM_015893	1.79
nuclear receptor subfamily 4, group A, member 3	U12767	1.79
T-box 2	AW303460	1.79
Consensus includes gb:AI952772 /FEA=EST /DB_XREF=gi:5745082 /DB_XREF=est:wr50d05.x1 /CLONE=IMAGE:2491113 /UG=Hs.300865 Homo sapiens clone BCSynL38 immunoglobulin h	AI952772	1.79
5-hydroxytryptamine (serotonin) receptor 2A	NM_000621	1.79
hypothetical protein FLJ23168	NM_025055	1.79
hypothetical protein PRO1446	NM_018601	1.79
glutamate decarboxylase 1 (brain, 67kD)	NM_013445	1.79
transcription factor ets	AF147782	1.79
forkhead box F1	NM_001451	1.79
hypothetical protein PRO0800	NM_018592	1.79
pre-B-cell leukemia transcription factor 2	BC003111	1.79
adrenergic, alpha-1A-, receptor	AF013261	1.79
short stature homeobox 2	AF022654	1.79
tachykinin receptor 1	M81797	1.79
Rev/Rex activation domain binding protein-related	AF015042	1.79
solute carrier family 2 (facilitated glucose transporter), member 3	NM_006931	1.79
hypothetical protein	AK023069	1.79
glutamate receptor, ionotropic, N-methyl D-aspartate 1	AF015731	1.79
Consensus includes gb:T87225 /FEA=EST /DB_XREF=gi:715577 /DB_XREF=est:yc81f06.s1 /CLONE=IMAGE:22392 /UG=Hs.13223 Homo sapiens mRNA full length insert cDNA clone EUROI	T87225	1.79
G protein-coupled receptor 22	NM_005295	1.79
Rhesus blood group-associated glycoprotein	AF178841	1.79
Consensus includes gb:AK025833.1 /DEF=Homo sapiens cDNA: FLJ22180 fis, clone HRC00936. /FEA=mRNA /DB_XREF=gi:10438467 /UG=Hs.287692 Homo sapiens cDNA: FLJ22180 fis, cl	AK025833	1.79
BAI1-associated protein 3	AB018277	1.79
fibrillin 2 (congenital contractural arachnodactyly)	X62009	1.79
nebulette	NM_016365	1.79
PDZ domain containing guanine nucleotide exchange factor(GEF)1	AV654984	1.79
collagen, type XI, alpha 1	NM_001854	1.79
cardiotrophin 1	NM_001330	1.79
epithelial membrane protein 1	NM_001423	1.79
synaptotagmin I	AV723167	1.79
Consensus includes gb:AU148024 /FEA=EST /DB_XREF=gi:11009545 /DB_XREF=est:AU148024 /CLONE=MAMMA1002392 /UG=Hs.302075 Homo sapiens cDNA FLJ12365 fis, clone MAMM	AU148024	1.79
cell division cycle 2-like 2	M37712	1.79
glutamate receptor, ionotropic, N-methyl D-aspartate 1	D13515	1.78
hypothetical protein FLJ11113	NM_018328	1.78
piccolo (presynaptic cytomatrix protein)	AC004082	1.78
son of sevenless (Drosophilia) homolog 2	BF692958	1.78
rabaptin-5	AF098638	1.78
KIAA1036 protein	NM_014909	1.78
stoned B/TFIIA-alpha/beta-like factor	NM_006873	1.78
SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily a, member 1	M88163	1.78
laminin, gamma 1 (formerly LAMB2)	I03202	1.78
D site of albumin promoter (albumin D-box) binding protein	D28468	1.78
erythropoietin receptor	X97671	1.78
zinc finger protein 222	NM_013360	1.78
gb:AF130057.1 /DEF=Homo sapiens clone FLB5539 PRO1454 mRNA, complete cds. /FEA=mRNA /PROD=PRO1454 /DB_XREF=gi:11493420 /UG=Hs.323288 Homo sapiens clone FLB5539	AF130057	1.78
platelet-derived growth factor receptor, beta polypeptide	NM_002609	1.78

DMC1 dosage suppressor of mck1 homolog, meiosis-specific homologous recombination (yeast)	D63882	1.78
endothelin 1	NM_001955	1.78
NBR2	NM_005821	1.78
Consensus includes gb:AK025029.1 /DEF=Homo sapiens cDNA: FLJ21376 fis, clone COL03231. /FEA=mRNA /DB_XREF=gi:10437464 /UG=Hs.306748 Homo sapiens cDNA: FLJ21376 fis, cl	AK025029	1.78
catenin (cadherin-associated protein), alpha 2	NM_004389	1.78
spondin 1, (f-spondin) extracellular matrix protein	AI885290	1.78
phosphodiesterase 1C, calmodulin-dependent (70kD)	NM_005020	1.78
radixin	NM_002906	1.78
calmodulin 2 (phosphorylase kinase, delta)	BC000454	1.78
Consensus includes gb:AV717062 /FEA=EST /DB_XREF=gi:10814214 /DB_XREF=est:AV717062 /CLONE=DCBCEG06 /UG=Hs.274553 Homo sapiens cDNA FLJ10906 fis, clone OVARC10000	AV717062	1.78
uncharacterized hematopoietic stem/progenitor cells protein MDS029	NM_018464	1.78
stathmin-like 4	NM_030795	1.78
hypothetical protein DKFZp761G1913	AL136557	1.78
SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily e, member 1	AL137010	1.78
RAB33B, member RAS oncogene family	NM_031296	1.78
neuropilin 2	AF280546	1.78
signal transducing adaptor molecule (SH3 domain and ITAM motif) 2	NM_005843	1.78
zinc finger protein 193	NM_006299	1.78
gastrulation brain homeo box 2	AF118452	1.78
hypothetical protein FLJ10460	NM_018097	1.78
hairy homolog (Drosophila)	NM_005524	1.78
KIAA0318 protein	AB002316	1.78
gb:U61167.1 /DEF=Human SH3 domain-containing protein SH3P18 mRNA, complete cds. /FEA=mRNA /PROD=SH3 domain-containing protein SH3P18 /DB_XREF=gi:1438934 /UG=Hs.330	U61167	1.78
mitogen-activated protein kinase 8 interacting protein 2	NM_012324	1.78
mitochondrial ribosomal protein L9	BE792072	1.78
lengsin	NM_016571	1.78
cofactor required for Sp1 transcriptional activation, subunit 9 (33kD)	BC005250	1.78
hypothetical protein PRO2822	NM_018539	1.78
topoisomerase (DNA) I	AW025108	1.78
calcium binding protein P22	NM_007236	1.78
KIAA0871 protein	BF224071	1.78
Consensus includes gb:BF940043 /FEA=EST /DB_XREF=gi:12357363 /DB_XREF=est:nac66f12.x1 /CLONE=IMAGE:3439271 /UG=Hs.62041 nidogen (enactin) /FL=gb:M30269.1 gb:NM_0(	BF940043	1.78
Cdc42 guanine exchange factor (GEF) 9	AI625739	1.78
protein kinase, AMP-activated, gamma 2 non-catalytic subunit	NM_016203	1.78
hypocretin (orexin) receptor 1	NM_001525	1.78
carbonic anhydrase VA, mitochondrial	NM_001739	1.78
sphingomyelin phosphodiesterase 1, acid lysosomal (acid sphingomyelinase)	M59917	1.78
Consensus includes gb:W28849 /FEA=EST /DB_XREF=gi:1308815 /DB_XREF=est:52g6 /UG=Hs.33862 ESTs	W28849	1.78
hypothetical protein FLJ20055	NM_017643	1.78
chromosome 5 open reading frame 4	NM_016348	1.78
hypothetical protein FLJ23497	NM_025089	1.78
Consensus includes gb:AL050043.1 /DEF=Homo sapiens mRNA; cDNA DKFZp566M0524 (from clone DKFZp566M0524). /FEA=mRNA /DB_XREF=gi:4884285 /UG=Hs.306310 Homo sapiens	AL050043	1.78
interleukin 6 receptor	NM_000565	1.78
Consensus includes gb:AL110248.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434N021 (from clone DKFZp434N021). /FEA=mRNA /DB_XREF=gi:5817198 /UG=Hs.208769 Homo sapiens m	AL110248	1.78
phosphodiesterase 1B, calmodulin-dependent	NM_000924	1.77
aldehyde dehydrogenase 1 family, member A2	AB015228	1.77
ret finger protein-like 3	NM_006604	1.77
EGF-like repeats and discoidin I-like domains 3	BF058643	1.77
ankyrin 2, neuronal	NM_001148	1.77
Consensus includes gb:AL117546.1 /DEF=Homo sapiens mRNA; cDNA DKFZp586C1923 (from clone DKFZp586C1923). /FEA=mRNA /DB_XREF=gi:5912079 /UG=Hs.241429 Homo sapiens	AL117546	1.77
A kinase (PRKA) anchor protein 5	NM_004857	1.77
leukocyte immunoglobulin-like receptor, subfamily A (without TM domain), member 3	NM_006865	1.77
tryptase beta 2	AF099143	1.77
Consensus includes gb:AK026856.1 /DEF=Homo sapiens cDNA: FLJ23203 fis, clone ADKA02487. /FEA=mRNA /DB_XREF=gi:10439812 /UG=Hs.306889 Homo sapiens cDNA: FLJ23203 fis,	AK026856	1.77
adaptor-related protein complex 1, sigma 1 subunit	BC003561	1.77
Consensus includes gb:R43279 /FEA=EST /DB_XREF=gi:821386 /DB_XREF=est:yg17c12.s1 /CLONE=IMAGE:32731 /UG=Hs.22574 ESTs	R43279	1.77
hypothetical protein FLJ20716	NM_017938	1.77
mitogen-activated protein kinase 8	U35004	1.77
Consensus includes gb:AF070543.1 /DEF=Homo sapiens clone 24740 mRNA sequence. /FEA=mRNA /DB_XREF=gi:3387903 /UG=Hs.155915 Homo sapiens clone 24740 mRNA sequence	AF070543	1.77
sialic acid binding Ig-like lectin 8	AF287892	1.77
Consensus includes gb:AK022686.1 /DEF=Homo sapiens cDNA FLJ12624 fis, clone NT2RM4001754. /FEA=mRNA /DB_XREF=gi:10434225 /UG=Hs.306639 Homo sapiens cDNA FLJ12624 f	AK022686	1.77
hypothetical protein FLJ13059	AK023121	1.77
Consensus includes gb:AL356504 /DEF=Human DNA sequence from clone RP1-14N1 on chromosome 1q21.1-21.3 Contains ESTs, GSSs and STSs. Contains the FLG gene for profilaggrin and	AL356504	1.77

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T cell receptor beta variable 17	U03115	1.77
amine oxidase, copper containing 3 (vascular adhesion protein 1)	NM_003734	1.77
ribosomal protein L18	AI669349	1.77
sodium channel, voltage-gated, type I, alpha polypeptide	AF225985	1.77
protein phosphatase 2, regulatory subunit B (B56), gamma isoform	AW772123	1.77
solute carrier family 10 (sodium/bile acid cotransporter family), member 2	NM_000452	1.77
deiodinase, iodothyronine, type I	NM_000792	1.77
integral membrane protein 2A	AL021786	1.77
methyl-CpG binding domain protein 2	AI827820	1.77
Consensus includes gb:AK023918.1 /DEF=Homo sapiens cDNA FLJ13856 fis, clone THYRO1000988. /FEA=mRNA /DB_XREF=gi:10436003 /UG=Hs.288489 Homo sapiens cDNA FLJ13856 fi	AK023918	1.77
sarcoglycan, gamma (35kD dystrophin-associated glycoprotein)	NM_000231	1.77
golgin-67	AW006438	1.77
anaplastic lymphoma kinase (Ki-1)	NM_004304	1.77
hypothetical protein PRO2268	NM_018520	1.77
hypothetical protein FLJ13942	NM_024581	1.77
glycine receptor, alpha 3	NM_006529	1.77
medium-chain acyl-CoA synthetase	AC003034	1.77
apolipoprotein C-I	NM_001645	1.77
KIAA1110 protein	AB029033	1.77
Consensus includes gb:AW405975 /FEA=EST /DB_XREF=gi:6925032 /DB_XREF=est:UI-HF-BL0-acv-a-05-0-UI.r1 /CLONE=IMAGE:3060368 /UG=Hs.247721 Homo sapiens clone mcg53-54 i	AW405975	1.77
carboxylesterase 3 (brain)	NM_012122	1.77
keratin 6A	J00269	1.77
solute carrier family 7 (cationic amino acid transporter, y+ system), member 8	AL365343	1.77
phosphodiesterase 4A, cAMP-specific (phosphodiesterase E2 dunce homolog, Drosophila)	L20965	1.77
CD6 antigen	U66146	1.77
Consensus includes gb:AF055011.1 /DEF=Homo sapiens clone 24587 mRNA sequence. /FEA=mRNA /DB_XREF=gi:3005734 /UG=Hs.137173 Homo sapiens clone 24587 mRNA sequence	AF055011	1.77
Consensus includes gb:AU158606 /FEA=EST /DB_XREF=gi:11020127 /DB_XREF=est:AU158606 /CLONE=PLACE3000406 /UG=Hs.287636 Homo sapiens cDNA FLJ14319 fis, clone PLACE3(	AU158606	1.77
Consensus includes gb:AL080086.1 /DEF=Homo sapiens mRNA; cDNA DKFZp564J1762 (from clone DKFZp564J1762). /FEA=mRNA /DB_XREF=gi:5262502 /UG=Hs.274538 Homo sapiens n	AL080086	1.77
inositol 1,4,5-trisphosphate 3-kinase A	NM_002220	1.77
Consensus includes gb:BC002465.1 /DEF=Homo sapiens, clone IMAGE:3344506, mRNA, partial cds. /FEA=mRNA /PROD=Unknown (protein for IMAGE:3344506) /DB_XREF=gi:12803298 /L	BC002465	1.77
S100 calcium binding protein A8 (calgranulin A)	AW238654	1.77
Consensus includes gb:AU157200 /FEA=EST /DB_XREF=gi:11018721 /DB_XREF=est:AU157200 /CLONE=PLACE1006617 /UG=Hs.301390 Homo sapiens cDNA FLJ13538 fis, clone PLACE1(	AU157200	1.77
GCN5 general control of amino-acid synthesis 5-like 1 (yeast)	NM_001487	1.77
testican 3	NM_016950	1.77
DKFZP586A0522 protein	AF113007	1.76
activin A receptor, type IB	NM_020327	1.76
epidermal growth factor receptor (erythroblastic leukemia viral (v-erb-b) oncogene homolog, avian)	U95089	1.76
Lsm1 protein	NM_014462	1.76
1,2-alpha-mannosidase IC	NM_020379	1.76
pre-mRNA processing factor 31 homolog (yeast)	AI334332	1.76
Consensus includes gb:AK022235.1 /DEF=Homo sapiens cDNA FLJ12173 fis, clone MAMMA1000696. /FEA=mRNA /DB_XREF=gi:10433588 /UG=Hs.306630 Homo sapiens cDNA FLJ12173 i	AK022235	1.76
protocadherin beta 11	NM_018931	1.76
T-cell leukemia/lymphoma 6	NM_012468	1.76
calpain 5	NM_004055	1.76
Consensus includes gb:AA555096 /FEA=EST /DB_XREF=gi:2325635 /DB_XREF=est:nj40e01.s1 /CLONE=IMAGE:994968 /UG=Hs.26295 Homo sapiens mRNA; cDNA DKFZp586D1122 (from	AA555096	1.76
solute carrier family 6 (neurotransmitter transporter, creatine), member 8	AW276522	1.76
lethal (3) malignant brain tumor l(3)mbt protein (Drosophila) homolog	AW445040	1.76
UV-B repressed sequence, HUR 7	BE148534	1.76
postmeiotic segregation increased 2-like 6	D38503	1.76
sirtuin silent mating type information regulation 2 homolog 2 (S. cerevisiae)	AI090841	1.76
protein tyrosine phosphatase, non-receptor type substrate 1	AL049634	1.76
follicular lymphoma variant translocation 1	NM_002035	1.76
myosin X	NM_012334	1.76
epithelial V-like antigen 1	NM_005797	1.76
ubiquitin-conjugating enzyme E2 variant 2	AL031291	1.76
gb:NM_018634.1 /DEF=Homo sapiens hypothetical protein PRO2893 (PRO2893), mRNA. /FEA=mRNA /GEN=PRO2893 /PROD=hypothetical protein PRO2893 /DB_XREF=gi:8924214 /UG=	NM_018634	1.76
Consensus includes gb:BF968134 /FEA=EST /DB_XREF=gi:12335349 /DB_XREF=est:602269121F1 /CLONE=IMAGE:4357349 /UG=Hs.250723 FK506 binding protein 12-rapamycin associa	BF968134	1.76
aquaporin 6, kidney specific	AL137716	1.76
proteinase 3 (serine proteinase, neutrophil, Wegener granulomatosis autoantigen)	NM_002777	1.76
gb:AF119889.1 /DEF=Homo sapiens PRO2667 mRNA, complete cds. /FEA=mRNA /PROD=PRO2667 /DB_XREF=gi:7770214 /UG=Hs.321170 Homo sapiens PRO2667 mRNA, complete cds /	AF119889	1.76
STAT induced STAT inhibitor-2	AB004903	1.76
membrane-spanning 4-domains, subfamily A, member 4A	NM_024021	1.76
Consensus includes gb:AW974995 /FEA=EST /DB_XREF=gi:8166198 /DB_XREF=est:EST387100 /UG=Hs.292787 ESTs	AW974995	1.76

tumor necrosis factor (ligand) superfamily, member 9	NM_003811	1.76
aggrecan 1 (chondroitin sulfate proteoglycan 1, large aggregating proteoglycan, antigen identified by monoclonal antibody A0122)	X17406	1.76
solute carrier family 6 (neurotransmitter transporter, creatine), member 10	U41163	1.76
G protein-coupled receptor 58	NM_014626	1.76
G protein-coupled receptor 4	NM_005282	1.76
adrenergic, alpha-1A-, receptor	D32201	1.76
ATP-binding cassette, sub-family B (MDR/TAP), member 11	NM_003742	1.76
Consensus includes gb:AK025180.1 /DEF=Homo sapiens cDNA: FLJ21527 fis, clone COL05961. /FEA=mRNA /DB_XREF=gi:10437643 /UG=Hs.306779 Homo sapiens cDNA: FLJ21527 fis, cl	AK025180	1.76
golgi phosphoprotein 1	AI636775	1.76
retinoblastoma binding protein 2	AF007135	1.76
LIM domain only 4	AI824831	1.76
cellular retinoic acid binding protein 2	NM_001878	1.76
Consensus includes gb:AI886482 /FEA=EST /DB_XREF=gi:5591646 /DB_XREF=est:wn39e07.x1 /CLONE=IMAGE:2447844 /UG=Hs.236750 ESTs, Highly similar to serine kinase SRPK2 H.sa	AI886482	1.76
immunoglobulin superfamily, member 4	AL519710	1.76
Consensus includes gb:BE326381 /FEA=EST /DB_XREF=gi:9200157 /DB_XREF=est:hw02e08.x1 /CLONE=IMAGE:3181766 /UG=Hs.18724 Homo sapiens mRNA; cDNA DKFZp564F093 (fron	BE326381	1.76
Consensus includes gb:BF110551 /FEA=EST /DB_XREF=gi:10940241 /DB_XREF=est:7n38h09.x1 /CLONE=IMAGE:3567208 /UG=Hs.253600 ESTs, Highly similar to S50128 glutamate rece	BF110551	1.76
Consensus includes gb:AK000122.1 /DEF=Homo sapiens cDNA FLJ20115 fis, clone COL05594. /FEA=mRNA /DB_XREF=gi:7020002 /UG=Hs.306382 Homo sapiens cDNA FLJ20115 fis, clon	AK000122	1.76
potassium voltage-gated channel, KQT-like subfamily, member 1	U89364	1.76
transducin-like enhancer of split 1 (E(sp1) homolog, Drosophila)	AI758763	1.76
Sm protein F	NM_007080	1.76
SH3-domain GRB2-like 3	AF036272	1.76
solute carrier family 7 (cationic amino acid transporter, y+ system), member 4	AW135332	1.76
hypothetical protein MGC3178	AI123471	1.76
karyopherin alpha 1 (importin alpha 5)	NM_002264	1.76
zinc finger protein 236	NM_007345	1.76
lymphocyte antigen 117	AF031137	1.75
tumor necrosis factor (ligand) superfamily, member 6	D38122	1.75
Consensus includes gb:AA479278 /FEA=EST /DB_XREF=gi:2207834 /DB_XREF=est:zv21b07.s1 /CLONE=IMAGE:754261 /UG=Hs.9295 elastin (supravalvular aortic stenosis, Williams-Beure	AA479278	1.75
H4 histone family, member A	NM_003538	1.75
chromosome 1 open reading frame 24	AF288391	1.75
aldehyde dehydrogenase 7 family, member A1	BC002515	1.75
tetraspan 5	AA748177	1.75
Consensus includes gb:AW204383 /FEA=EST /DB_XREF=gi:6503855 /DB_XREF=est:UI-H-BI1-adv-c-03-0-UI.s1 /CLONE=IMAGE:2717957 /UG=Hs.312723 ESTs, Highly similar to F22162 1	AW204383	1.75
major histocompatibility complex, class I-like sequence	AF010447	1.75
bombesin-like receptor 3	Z97632	1.75
KIAA0138 gene product	D50928	1.75
UDP glycosyltransferase 1 family, polypeptide A8	NM_019076	1.75
protease, serine, 7 (enterokinase)	NM_002772	1.75
DNA segment on chromosome X (unique) 9879 expressed sequence	AA034498	1.75
prothymosin, alpha (gene sequence 28)	BF686442	1.75
hypothetical protein FLJ11292	AK023539	1.75
hypothetical protein PRO1257	NM_018578	1.75
hypothetical protein PRO2289	NM_018625	1.75
coatomer protein complex, subunit beta	NM_016451	1.75
purine-rich element binding protein G	NM_013357	1.75
DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 6 (RNA helicase, 54kD)	NM_004397	1.75
sperm adhesion molecule 1 (PH-20 hyaluronidase, zona pellucida binding)	S67798	1.75
SH3-domain binding protein 4	AF015043	1.75
poliovirus receptor	NM_006505	1.75
cadherin 18, type 2	NM_004934	1.75
vanilloid receptor-related osmotically activated channel; OTRPC4 protein	NM_021625	1.75
Consensus includes gb:BG389073 /FEA=EST /DB_XREF=gi:13282519 /DB_XREF=est:602414868F1 /CLONE=IMAGE:4523201 /UG=Hs.151010 ESTs	BG389073	1.75
programmed cell death 2	AA764988	1.75
hypothetical protein FLJ11029	NM_018304	1.75
pp21 homolog	NM_016303	1.75
annexin A2 pseudogene 1	M62896	1.75
hypothetical protein FLJ20241	AK023399	1.75
integrin, alpha 5 (fibronectin receptor, alpha polypeptide)	NM_002205	1.75
Consensus includes gb:AF035594.1 /DEF=Homo sapiens protein kinase C-alpha mRNA, partial 3 UTR. /FEA=mRNA /DB_XREF=gi:3168857 /UG=Hs.279856 Homo sapiens protein kinase C-a	AF035594	1.75
Consensus includes gb:AW188214 /FEA=EST /DB_XREF=gi:6462650 /DB_XREF=est:xj93g09.x1 /CLONE=IMAGE:2664832 /UG=Hs.250389 ESTs, Weakly similar to A59252 myosin heavy ch	AW188214	1.75
RAD51-like 1 (S. cerevisiae)	Y15571	1.75
KIAA1305 protein	NM_025081	1.75
hypothetical protein FLJ13710	NM_024817	1.75

sperm acrosome associated 1	NM_030960	1.75
glial cells missing homolog b (Drosophila)	NM_004752	1.75
peptidylprolyl isomerase C (cyclophilin C)	BE962749	1.75
KIAA0443 gene product	NM_014710	1.75
purine-rich element binding protein A	BE222739	1.75
phosphodiesterase 1C, calmodulin-dependent (70kD)	U40372	1.75
echinoderm microtubule-associated protein-like	AI825937	1.75
hypothetical protein FLJ11036	NM_018306	1.75
Consensus includes gb:AK021514.1 /DEF=Homo sapiens cDNA FLJ11452 fis, clone HEMBA1001435. /FEA=mRNA /DB_XREF=gi:10432710 /UG=Hs.148598 Homo sapiens cDNA FLJ11452 f	AK021514	1.75
pancreatic polypeptide 2	NM_021092	1.75
melanoma cell adhesion molecule	M29277	1.75
Consensus includes gb:AK024158.1 /DEF=Homo sapiens cDNA FLJ14096 fis, clone MAMMA1000752. /FEA=mRNA /DB_XREF=gi:10436469 /UG=Hs.303289 Homo sapiens cDNA FLJ14096 f	AK024158	1.75
lysosomal-associated membrane protein 3	NM_014398	1.75
recombination activating gene 1	NM_000448	1.75
prospero-related homeobox 1	NM_002763	1.75
alpha-methylacyl-CoA racemase	AK022765	1.74
Consensus includes gb:X16454 /DEF=Human gene for carcinoembryonic antigen subdomains A and B /FEA=CDS /DB_XREF=gi:29852 /UG=Hs.248025 Human gene for carcinoembryonic an	X16454	1.74
zinc finger protein 146	NM_007145	1.74
T-cell acute lymphocytic leukemia 1	X51990	1.74
muscle, skeletal, receptor tyrosine kinase	NM_005592	1.74
Consensus includes gb:AL133075.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434M0317 (from clone DKFZp434M0317). /FEA=mRNA /DB_XREF=gi:6453518 /UG=Hs.274560 Homo sapiens	AL133075	1.74
Apobec-1 complementation factor; APOBEC-1 stimulating protein	NM_014576	1.74
gb:BC002778.1 /DEF=Homo sapiens, Similar to myosin light chain 2, precursor lymphocyte-specific, clone MGC:3479, mRNA, complete cds. /FEA=mRNA /PROD=Similar to myosin light chai	BC002778	1.74
mannose-6-phosphate receptor (cation dependent)	AI583537	1.74
Consensus includes gb:AK026439.1 /DEF=Homo sapiens cDNA: FLJ22786 fis, clone KAIA2150. /FEA=mRNA /DB_XREF=gi:10439304 /UG=Hs.306848 Homo sapiens cDNA: FLJ22786 fis, clo	AK026439	1.74
inhibin, beta B (activin AB beta polypeptide)	NM_002193	1.74
transmembrane 4 superfamily member (tetraspan NET-2)	NM_012338	1.74
T-cell receptor interacting molecule	AI240085	1.74
pipecolic acid oxidase	AF136970	1.74
aldehyde dehydrogenase 3 family, member B2	NM_000695	1.74
chondroitin sulfate proteoglycan 4 (melanoma-associated)	NM_001897	1.74
carboxypeptidase B2 (plasma, carboxypeptidase U)	NM_016413	1.74
coagulation factor II (thrombin) receptor-like 1	NM_005242	1.74
KIAA0675 gene product	NM_014648	1.74
cystatin A (stefin A)	NM_005213	1.74
neurexin 1	AB035356	1.74
vacuolar protein sorting 16 (yeast)	AA779495	1.74
Consensus includes gb:AW295367 /FEA=EST /DB_XREF=gi:6702003 /DB_XREF=est:UI-H-BI2-ahv-e-04-0-UI.s1 /CLONE=IMAGE:2728039 /UG=Hs.196585 ESTs, Weakly similar to Pro-Pol-d	AW295367	1.74
cathepsin B	W47179	1.74
olfactory receptor, family 10, subfamily H, member 3	NM_013938	1.74
Consensus includes gb:U82306.1 /DEF=Homo sapiens unknown protein mRNA, partial cds. /FEA=mRNA /PROD=unknown /DB_XREF=gi:1938333 /UG=Hs.123081 Homo sapiens unknown p	U82306	1.74
left-right determination, factor B	NM_020997	1.74
Consensus includes gb:AL079289.1 /DEF=Homo sapiens mRNA full length insert cDNA clone EUROIMAGE 35971. /FEA=mRNA /DB_XREF=gi:5102748 /UG=Hs.137154 Homo sapiens mRNA	AL079289	1.74
Consensus includes gb:AI251844.1 /DEF=Homo sapiens partial mRNA for MOZCBP chimeric transcript type II. /FEA=mRNA /GEN=MOZCBP /PROD=MOZCBP /DB_XREF=gi:9368433 /UG=Hs.	AI251844	1.74
angiotensinogen (serine (or cysteine) proteinase inhibitor, clade A (alpha-1 antiproteinase, antitrypsin), member 8)	NM_000029	1.74
hypothetical protein FLJ14031	AK024093	1.74
fibroblast growth factor receptor 2 (bacteria-expressed kinase, keratinocyte growth factor receptor, craniofacial dysostosis 1, Crouzon syndrome, Pfeiffer syndrome, Jackson-Weiss syndrome)	AB030075	1.74
tripeptidyl peptidase II	NM_003291	1.74
transcription factor binding to IGHM enhancer 3	NM_006521	1.74
ATP-binding cassette, sub-family A (ABC1), member 1	AK024328	1.74
solute carrier family 22 (organic cation transporter), member 5	NM_003060	1.74
hypothetical protein FLJ12457	NM_024674	1.74
cadherin 19, type 2	NM_021153	1.74
retinoic acid receptor, beta	NM_016152	1.74
neurotrophic tyrosine kinase, receptor, type 1	X04201	1.74
prolactin receptor	S78505	1.74
Consensus includes gb:AK024860.1 /DEF=Homo sapiens cDNA: FLJ21207 fis, clone COL00362. /FEA=mRNA /DB_XREF=gi:10437268 /UG=Hs.306708 Homo sapiens cDNA: FLJ21207 fis, cl	AK024860	1.74
hypothetical protein FLJ10209	NM_018026	1.74
GAP-associated tyrosine phosphoprotein p62 (Sam68)	AW592227	1.74
hypothetical protein FLJ10488	BC003541	1.74
rab3 GTPase-activating protein, non-catalytic subunit (150kD)	AF255648	1.74
RNA binding motif protein 5	AF107493	1.74
Consensus includes gb:R38712 /FEA=EST /DB_XREF=gi:796168 /DB_XREF=est:yd03b02.s1 /CLONE=IMAGE:24533 /UG=Hs.155959 Homo sapiens clone 24533 mRNA sequence	R38712	1.74

ubiquitin protein ligase E3A (human papilloma virus E6-associated protein, Angelman syndrome)	U84404	1.74
CS box-containing WD protein	BG260658	1.74
hypocretin (orexin) receptor 2	NM_001526	1.74
cytokeratin 2	NM_015848	1.74
hypothetical protein FLJ22601	NM_024822	1.74
chromosome 4 open reading frame 5	NM_016599	1.74
annexin A5	NM_001154	1.74
growth differentiation factor 2	NM_016204	1.74
hypothetical protein FLJ13448	NM_025147	1.74
forkhead box D2	NM_004474	1.74
glycerol-3-phosphate dehydrogenase 1 (soluble)	NM_005276	1.74
hypothetical protein FLJ22031	NM_025074	1.74
insulin-like growth factor binding protein, acid labile subunit	AW338791	1.74
kallikrein 14	NM_022046	1.74
Consensus includes gb:AI275371 /DEF=Homo sapiens partial IGVH3 gene for immunoglobulin heavy chain V region, case 1, clone 16 /FEA=CDS /DB_XREF=gi:7573002 /UG=Hs.272356 Ho	AI275371	1.74
Consensus includes gb:U82306.1 /DEF=Homo sapiens unknown protein mRNA, partial cds. /FEA=mRNA /PROD=unknown /DB_XREF=gi:1938333 /UG=Hs.123081 Homo sapiens unknown p	U82306	1.74
poliovirus receptor-related 2 (herpesvirus entry mediator B)	NM_002856	1.73
L38424 B subtilis dapB, jojF, joj6 genes corresponding to nucleotides 1358-3197 of L38424 (-5, -M, -3 represent transcript regions 5 prime, Middle, and 3 prime respectively)	L38424	1.73
four and a half LIM domains 1	AF220153	1.73
PERB11 family member in MHC class I region	NM_005844	1.73
hypothetical protein FLJ10035	NM_017974	1.73
CD1E antigen, e polypeptide	NM_030893	1.73
Consensus includes gb:AA443771 /FEA=EST /DB_XREF=gi:2156446 /DB_XREF=est:zw95f08.s1 /CLONE=IMAGE:784743 /UG=Hs.270138 ESTs	AA443771	1.73
discs, large (Drosophila) homolog 5	AB011155	1.73
Consensus includes gb:BC005092.1 /DEF=Homo sapiens, clone IMAGE:3843012, mRNA, partial cds. /FEA=mRNA /PROD=Unknown (protein for IMAGE:3843012) /DB_XREF=gi:13477248 /L	BC005092	1.73
gb:AF116715.1 /DEF=Homo sapiens PRO2829 mRNA, complete cds. /FEA=mRNA /PROD=PRO2829 /DB_XREF=gi:7959928 /UG=Hs.256256 Homo sapiens PRO2829 mRNA, complete cds /	AF116715	1.73
collagen, type I, alpha 1	NM_000088	1.73
golgi autoantigen, golgin subfamily b, macrogolgin (with transmembrane signal), 1	N53479	1.73
secreted protein of unknown function	BF440021	1.73
PRO0618 protein	NM_014133	1.73
otoraplin	NM_020157	1.73
leukocyte immunoglobulin-like receptor, subfamily B (with TM and ITIM domains), member 3	AF009644	1.73
caspase 5, apoptosis-related cysteine protease	NM_004347	1.73
hypothetical protein FLJ20761	NM_017954	1.73
MSPC048 protein	NM_014148	1.73
potassium inwardly-rectifying channel, subfamily J, member 5	L47208	1.73
microsomal NAD+-dependent retinol dehydrogenase 4	AF086735	1.73
solute carrier family 30 (zinc transporter), member 4	NM_013309	1.73
RAB2, member RAS oncogene family	AW301641	1.73
matrix metalloproteinase 16 (membrane-inserted)	NM_022564	1.73
carboxypeptidase, vitellogenic-like	NM_031311	1.73
zinc finger protein 141 (clone pHZ-44)	NM_003441	1.73
Consensus includes gb:AK022686.1 /DEF=Homo sapiens cDNA FLJ12624 fis, clone NT2RM4001754. /FEA=mRNA /DB_XREF=gi:10434225 /UG=Hs.306639 Homo sapiens cDNA FLJ12624 fi	AK022686	1.73
Consensus includes gb:AA573805 /FEA=EST /DB_XREF=gi:2348320 /DB_XREF=est:nk07g11.s1 /CLONE=IMAGE:1012868 /UG=Hs.152939 Homo sapiens clone 24630 mRNA sequence	AA573805	1.73
deleted in bladder cancer chromosome region candidate 1	NM_014618	1.73
lipase protein	BC001698	1.73
KIAA1046 protein	NM_014928	1.73
hypothetical protein FLJ20337	AL096712	1.73
prostate androgen-regulated transcript 1	NM_016590	1.73
calmodulin-like skin protein	NM_017422	1.73
DKFZP586P0123 protein	AL080220	1.73
KIAA0485 protein	AI122905	1.73
solute carrier family 26, member 3	NM_000111	1.73
Consensus includes gb:AA017721 /FEA=EST /DB_XREF=gi:1479910 /DB_XREF=est:ze39f11.s1 /CLONE=IMAGE:361389 /UG=Hs.49117 Homo sapiens mRNA; cDNA DKFZp564N1662 (from	AA017721	1.73
rat regenerating islet-derived-like, human homolog (pancreatic stone protein-like, pancreatic thread protein-like)	NM_006508	1.73
SRY (sex determining region Y)-box 9 (campomelic dysplasia, autosomal sex-reversal)	NM_000346	1.73
RuvB-like 1 (E. coli)	M34421	1.73
Consensus includes gb:AL137292.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434L1016 (from clone DKFZp434L1016). /FEA=mRNA /DB_XREF=gi:6807752 /UG=Hs.274573 Homo sapiens	AL137292	1.73
Consensus includes gb:K00627.1 /DEF=human kpni repeat mrna (cdna clone pcd-kpni-8), 3 end. /FEA=mRNA /DB_XREF=gi:337653 /UG=Hs.203776 Human kpni repeat mrna (cdna clone	K00627	1.73
hypothetical protein FLJ13166	NM_025003	1.73
Consensus includes gb:AW149844 /FEA=EST /DB_XREF=gi:6197740 /DB_XREF=est:xf42h10.x1 /CLONE=IMAGE:2620771 /UG=Hs.167140 ESTs, Moderately similar to CIKG_HUMAN VOLTA	AW149844	1.73
cyclin-dependent kinase 6	AI738463	1.73
secreted frizzled-related protein 5	NM_003015	1.73

Epstein-Barr virus induced gene 2 (lymphocyte-specific G protein-coupled receptor)	NM_004951	1.73
hypothetical protein FLJ11561	NM_024651	1.73
hypothetical protein FLJ20637	NM_017912	1.73
tumor necrosis factor, alpha-induced protein 6	NM_007115	1.73
Consensus includes gb:AV691491 /FEA=EST /DB_XREF=gi:10293354 /DB_XREF=est:AV691491 /CLONE=GKCEEB10 /UG=Hs.85335 Homo sapiens mRNA; cDNA DKFZp564D1462 (from clor	AV691491	1.73
KIAA0478 gene product	NM_014870	1.73
mannosyl (alpha-1,3-)-glycoprotein beta-1,4-N-acetylglucosaminyltransferase, isoenzyme B	NM_014275	1.73
phosphodiesterase 4C, cAMP-specific (phosphodiesterase E1 dunce homolog, Drosophila)	NM_000923	1.73
solute carrier family 7 (cationic amino acid transporter, y+ system), member 8	AL365343	1.72
Consensus includes gb:AL035295.1 /DEF=H.sapiens gene from PAC 106H8, similar to Dynamin. /FEA=mRNA /PROD=hypothetical protein /DB_XREF=gi:4200217 /UG=Hs.56175 H.sapiens	AL035295	1.72
MUF1 protein	NM_006369	1.72
6 antigen 78	NM_001477	1.72
olfactory receptor, family 12, subfamily D, member 2	NM_013936	1.72
hypothetical protein FLJ23091	AA775681	1.72
potassium inwardly-rectifying channel,subfamily J, member 13	AI007557	1.72
hypothetical protein FLJ12526	NM_024787	1.72
Consensus includes gb:AI143919 /FEA=EST /DB_XREF=gi:3665728 /DB_XREF=est:qd99f12.x1 /CLONE=IMAGE:1737647 /UG=Hs.193783 Human DNA sequence from clone RP13-329D4 or	AI143919	1.72
glutaminase	AF158555	1.72
hypothetical protein FLJ13352	NM_024592	1.72
ADP-ribosylation factor 6	NM_001663	1.72
hypothetical protein PRO2007	NM_018613	1.72
muscle-specific beta 1 integrin binding protein	NM_014446	1.72
splicing factor, arginine/serine-rich 6	NM_006275	1.72
Consensus includes gb:AK025388.1 /DEF=Homo sapiens cDNA: FLJ21735 fis, clone COLF3350. /FEA=mRNA /DB_XREF=gi:10437892 /UG=Hs.306808 Homo sapiens cDNA: FLJ21735 fis, cl	AK025388	1.72
neuropeptide Y receptor Y2	U32500	1.72
armadillo repeat protein ALEX2	NM_014782	1.72
sphingosine-1-phosphate lyase 1	NM_003901	1.72
Consensus includes gb:AF005082.1 /DEF=Homo sapiens skin-specific protein (xp33) mRNA, partial cds. /FEA=mRNA /GEN=xp33 /PROD=skin-specific protein /DB_XREF=gi:2589191 /UG=	AF005082	1.72
uncharacterized hypothalamus protein HT011	BE565675	1.72
Consensus includes gb:AK002179.1 /DEF=Homo sapiens cDNA FLJ11317 fis, clone PLACE1010261, moderately similar to SEGREGATION DISTORTER PROTEIN. /FEA=mRNA /DB_XREF=gi:10	AK002179	1.72
integrin, beta 8	BC002630	1.72
Consensus includes gb:AV760769 /FEA=EST /DB_XREF=gi:10918617 /DB_XREF=est:AV760769 /CLONE=MDSBSE09 /UG=Hs.306324 Homo sapiens mRNA; cDNA DKFZp434G182 (from clor	AV760769	1.72
KIAA0414 protein	T90308	1.72
ATP-binding cassette, sub-family A (ABC1), member 12	AL080207	1.72
H4 histone family, member B	NM_003539	1.72
amine oxidase, copper containing 3 (vascular adhesion protein 1)	AF047485	1.72
Consensus includes gb:BE786164 /FEA=EST /DB_XREF=gi:10207362 /DB_XREF=est:601474273F1 /CLONE=IMAGE:3877146 /UG=Hs.80285 Homo sapiens mRNA; cDNA DKFZp586C1723	BE786164	1.72
Consensus includes gb:BE677757 /FEA=EST /DB_XREF=gi:10038372 /DB_XREF=est:7f59g09.x1 /CLONE=IMAGE:3299008 /UG=Hs.311561 ESTs, Weakly similar to CYA7_HUMAN ADENYLA	BE677757	1.72
glycine-N-acyltransferase	NM_005838	1.72
neuronal thread protein	AF010144	1.72
complement component 1, r subcomponent	AL573058	1.72
PRO1693 protein	AF118071	1.72
par-3 partitioning defective 3 homolog (C. elegans)	AF196186	1.72
hypothetical protein FLJ20668	NM_017923	1.72
olfactory receptor, family 3, subfamily A, member 3	NM_012373	1.72
DKFZP434M131 protein	AL080179	1.72
hypothetical protein FLJ20069	NM_017651	1.72
glutamate receptor, metabotropic 8	AC000099	1.72
Consensus includes gb:AL133018.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434F0327 (from clone DKFZp434F0327). /FEA=mRNA /DB_XREF=gi:6453498 /UG=Hs.270527 Homo sapiens	AL133018	1.72
chromosome 11 open reading frame 8	NM_001584	1.72
hypothetical protein dJ796I17.1	NM_025225	1.72
taste receptor, type 2, member 8	NM_023918	1.72
Consensus includes gb:AL355532 /DEF=Human DNA sequence from clone RP11-487F5 on chromosome 6 Contains the 3 part of the GRIK2 (glutamate receptor, ionotropic, kainate 2) gene,	AL355532	1.72
interleukin 6 receptor	S72848	1.72
transforming growth factor, beta 2	BF061658	1.72
lipin 2	U55968	1.72
sphingomyelin phosphodiesterase 1, acid lysosomal (acid sphingomyelinase)	M81780	1.72
astrotactin	AB006627	1.72
Consensus includes gb:X81636.1 /DEF=H.sapiens clathrin light chain a gene. /FEA=mRNA /DB_XREF=gi:704460 /UG=Hs.285688 H.sapiens clathrin light chain a gene	X81636	1.72
neuralized-like (Drosophila)	AF029729	1.72
Consensus includes gb:AL117447.1 /DEF=Homo sapiens mRNA; cDNA DKFZp586A0617 (from clone DKFZp586A0617). /FEA=mRNA /DB_XREF=gi:5911895 /UG=Hs.306342 Homo sapiens	AL117447	1.72
hydatidiform mole associated and imprinted	AF241534	1.72
sex comb on midleg-like 2 (Drosophila)	NM_006089	1.72

Consensus includes gb:AU144530 /FEA=EST /DB_XREF=gi:11006051 /DB_XREF=est:AU144530 /CLONE=HEMBA1002189 /UG=Hs.296654 Homo sapiens cDNA FLJ11511 fis, clone HEMBA	AU144530	1.72
basonuclin	NM_001717	1.72
FYN binding protein (FYB-120/130)	AF198052	1.72
carbohydrate (N-acetylglucosamine 6-O) sulfotransferase 7	AL022165	1.72
aggrecan 1 (chondroitin sulfate proteoglycan 1, large aggregating proteoglycan, antigen identified by monoclonal antibody A0122)	NM_001135	1.72
KIAA1232 protein	T62872	1.72
tissue inhibitor of metalloproteinase 4	NM_003256	1.72
uncharacterized gastric protein ZA52P	NM_020669	1.72
hepatitis delta antigen-interacting protein A	BE966372	1.72
interferon, alpha 4	NM_021068	1.72
Consensus includes gb:AK022006.1 /DEF=Homo sapiens cDNA FLJ11944 fis, clone HEMBB1000684. /FEA=mRNA /DB_XREF=gi:10433318 /UG=Hs.300771 Homo sapiens cDNA FLJ11944 f	AK022006	1.72
leukocyte cell-derived chemotaxin 2	NM_002302	1.72
vitiligo-associated protein VIT-1	NM_018693	1.72
chromosome 16 open reading frame 5	NM_013399	1.71
wingless-type MMTV integration site family, member 10B	NM_003394	1.71
KIAA1096 protein	BG261071	1.71
bridging integrator 1	BG034080	1.71
multifunctional polypeptide similar to SAICAR synthetase and AIR carboxylase	AU121975	1.71
hypothetical protein FLJ20041	NM_017636	1.71
KIAA1107 protein	AB029030	1.71
estrogen receptor 1	AJ073549	1.71
Consensus includes gb:AF254357.1 /DEF=Homo sapiens prostate-specific membrane antigen PSM mRNA, exon 6 alternative splice variant, partial cds. /FEA=mRNA /PROD=prostate-specific	AF254357	1.71
carbonyl reductase 1	BC002511	1.71
restin (Reed-Steinberg cell-expressed intermediate filament-associated protein)	M97501	1.71
Consensus includes gb:AA628440 /FEA=EST /DB_XREF=gi:2540827 /DB_XREF=est:af26f02.s1 /CLONE=IMAGE:1032795 /UG=Hs.83623 nuclear receptor subfamily 1, group I, member 3	AA628440	1.71
keratin, hair, basic, 3	NM_002282	1.71
azurocidin 1 (cationic antimicrobial protein 37)	NM_001700	1.71
Consensus includes gb:AL109809 /DEF=Human DNA sequence from clone RP4-673D20 on chromosome 20. Contains a PTPN5 (protein tyrosine phosphatase, non-receptor type substrate) f	AL109809	1.71
Consensus includes gb:AV720842 /FEA=EST /DB_XREF=gi:10817994 /DB_XREF=est:AV720842 /CLONE=GLCCOH05 /UG=Hs.282435 ESTs	AV720842	1.71
ankyrin 3, node of Ranvier (ankyrin G)	NM_001149	1.71
serine (or cysteine) proteinase inhibitor, clade A (alpha-1 antiproteinase, antitrypsin), member 1	NM_000295	1.71
follistatin-like 1	BC000055	1.71
GDNF family receptor alpha 2	NM_001495	1.71
gb:BC002821.1 /DEF=Homo sapiens, clone MGC:3518, mRNA, complete cds. /FEA=mRNA /PROD=Unknown (protein for MGC:3518) /DB_XREF=gi:12803948 /UG=Hs.306987 Homo sapiens	BC002821	1.71
chimaerin (chimaerin) 2	L29126	1.71
glycine amidinotransferase (L-arginine:glycine amidinotransferase)	NM_001482	1.71
gastric inhibitory polypeptide receptor	NM_000164	1.71
leptin receptor	U50748	1.71
Consensus includes gb:AL157490.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434F1011 (from clone DKFZp434F1011). /FEA=mRNA /DB_XREF=gi:7018535 /UG=Hs.274589 Homo sapiens	AL157490	1.71
KIAA0857 protein	AF334812	1.71
transcription factor 4	AK026674	1.71
gb:NM_003723.1 /DEF=Homo sapiens caspase 13, apoptosis-related cysteine protease (CASP13), mRNA. /FEA=mRNA /GEN=CASP13 /PROD=caspase 13 /DB_XREF=gi:4502570 /UG=Hs.	NM_003723	1.71
glycolipid transfer protein	NM_016433	1.71
laminin, alpha 3 (nicein (150kD), kalinin (165kD), BM600 (150kD), epilegrin)	NM_000227	1.71
erythrocyte protein band 4.1-like 4	NM_022140	1.71
homeodomain interacting protein kinase 2	NM_022740	1.71
chemokine (C-C motif) receptor-like 2	AF015524	1.71
Consensus includes gb:AL047879 /FEA=EST /DB_XREF=gi:4728067 /DB_XREF=est:DKFZp586N1222_s1 /CLONE=DKFZp586N1222 /UG=Hs.194251 ESTs, Weakly similar to ALU2_HUMAN	AL047879	1.71
fibroblast activation protein, alpha	U76833	1.71
heat shock 70kD protein 1B	NM_005346	1.71
hypothetical protein FLJ10261	NM_018043	1.71
RPA-binding trans-activator	NM_013368	1.71
hypothetical protein FLJ10687	NM_018178	1.71
Consensus includes gb:U92027.1 /DEF=Human clone 61501 defective mariner transposon Hsmar2 mRNA sequence. /FEA=mRNA /DB_XREF=gi:2052469 /UG=Hs.159211 Human clone 61	U92027	1.71
cholinergic receptor, nicotinic, alpha polypeptide 3	BC000513	1.71
midline 1 (Opitz/BBB syndrome)	AF041209	1.71
Consensus includes gb:AK024851.1 /DEF=Homo sapiens cDNA: FLJ21198 fis, clone COL00220. /FEA=mRNA /DB_XREF=gi:10437259 /UG=Hs.306707 Homo sapiens cDNA: FLJ21198 fis, cl	AK024851	1.71
Consensus includes gb:AL109671.1 /DEF=Homo sapiens mRNA full length insert cDNA clone EUROIMAGE 29222. /FEA=mRNA /DB_XREF=gi:5689802 /UG=Hs.140833 Homo sapiens mRNA	AL109671	1.71
troponin T1, skeletal, slow	AJ011712	1.71
gb:U35622.2 /DEF=Homo sapiens EWS proteinE1A enhancer binding protein chimera mRNA, complete cds. /FEA=mRNA /PROD=EWS proteinE1A enhancer binding proteinchimera /DB_XREF	U35622	1.71
hypothetical protein FLJ21079	NM_024576	1.71
parathyroid hormone-like hormone	BC005961	1.71
folate receptor 2 (fetal)	NM_000803	1.71

deoxyribonuclease II, lysosomal	AB004574	1.71
myristoylated alanine-rich protein kinase C substrate	NM_002356	1.71
hypothetical protein FLJ21963	NM_024560	1.71
H2A histone family, member N	NM_003514	1.71
brain specific protein	NM_016140	1.71
splicing factor 3a, subunit 2, 66kD	BC004434	1.71
adrenergic, beta-3-, receptor	X70812	1.71
pparl	NM_016566	1.71
Ig superfamily protein	NM_007268	1.71
GPI anchored molecule like protein	NM_002066	1.71
myristoylated alanine-rich protein kinase C substrate	M68956	1.71
tumor protein p73	NM_005427	1.71
steroid-5-alpha-reductase, alpha polypeptide 2 (3-oxo-5 alpha-steroid delta 4-dehydrogenase alpha 2)	NM_000348	1.71
collagen, type VI, alpha 1	M20776	1.71
signal transducer and activator of transcription 2, 113kD	S81491	1.71
gb:AF130051.1 /DEF=Homo sapiens clone FLB3535 PR00898 mRNA, complete cds. /FEA=mRNA /PROD=PR00898 /DB_XREF=gi:11493408 /UG=Hs.306960 Homo sapiens clone FLB3535	AF130051	1.71
Consensus includes gb:AL136179 /DEF=Human DNA sequence from clone RP3-322L4 on chromosome 6. Contains the SOX4 gene for SRY (sex determining region Y)-box 4, a pseudogene s	AL136179	1.71
KIAA0557 protein	AB011129	1.71
tumor stroma and activated macrophage protein DLM-1	NM_030776	1.71
KIAA1000 protein	BF062942	1.71
tumor necrosis factor (ligand) superfamily, member 13	AF114013	1.71
lysyl oxidase-like 1	NM_005576	1.71
Consensus includes gb:BE858194 /FEA=EST /DB_XREF=gi:10372818 /DB_XREF=est:7g19c05.x1 /CLONE=IMAGE:3306920 /UG=Hs.323053 Homo sapiens mRNA full length insert cDNA clc	BE858194	1.70
Consensus includes gb:AU144277 /FEA=EST /DB_XREF=gi:11005798 /DB_XREF=est:AU144277 /CLONE=HEMBA1001415 /UG=Hs.296525 Homo sapiens cDNA FLJ10062 fis, clone HEMBA	AU144277	1.70
ATPase, Ca++ transporting, plasma membrane 1	L14561	1.70
solute carrier family 17 (sodium-dependent inorganic phosphate cotransporter), member 7	H40895	1.70
Consensus includes gb:AL157398 /DEF=Human DNA sequence from clone RP11-56H7 on chromosome 10. Contains ESTs, STSs and GSSs. Contains the gene for the nebulette protein (NEBL.	AL157398	1.70
protein (peptidyl-prolyl cis/trans isomerase) NIMA-interacting 1-like	NM_006222	1.70
transcription factor AP-2 beta (activating enhancer binding protein 2 beta)	NM_003221	1.70
transportin-SR	AK022910	1.70
zinc finger protein 214	NM_013249	1.70
Ric-like, expressed in neurons (Drosophila)	NM_002930	1.70
integrin, alpha 2b (platelet glycoprotein IIb of IIb/IIIa complex, antigen CD41B)	I02963	1.70
inositol 1,4,5-triphosphate receptor, type 1	L38019	1.70
ectonucleoside triphosphate diphosphohydrolase 3	NM_001248	1.70
purinergic receptor P2X, ligand-gated ion channel, 4	NM_002560	1.70
CCR4 carbon catabolite repression 4-like (S. cerevisiae)	NM_012118	1.70
Consensus includes gb:AU159412 /FEA=EST /DB_XREF=gi:11020933 /DB_XREF=est:AU159412 /CLONE=THYRO1001602 /UG=Hs.296756 Homo sapiens cDNA FLJ14348 fis, clone THYRO'	AU159412	1.70
retinoic acid repressible protein	AF172066	1.70
hypothetical protein FLJ14075	NM_024894	1.70
carnitine O-octanoyltransferase	NM_021151	1.70
surfactant, pulmonary-associated protein B	I02761	1.70
protein kinase, cGMP-dependent, type II	NM_006259	1.70
LRB protein	NM_014020	1.70
KIAA1654 protein	AB051441	1.70
c-myc binding protein	D50692	1.70
Consensus includes gb:AL161956.1 /DEF=Homo sapiens mRNA; cDNA DKFZp761A17121 (from clone DKFZp761A17121). /FEA=mRNA /DB_XREF=gi:7328008 /UG=Hs.306492 Homo sapie	AL161956	1.70
potassium channel, subfamily K, member 7	NM_005714	1.70
Sp4 transcription factor	NM_003112	1.70
6 protein-coupled receptor 86	NM_023914	1.70
transporter 2, ATP-binding cassette, sub-family B (MDR/TAP)	NM_018833	1.70
defensin, alpha 4, corticostatin	NM_001925	1.70
hypothetical protein, estradiol-induced	AF250321	1.70
hypothetical protein FKS625	NM_022120	1.70
hypothetical protein FLJ23093	NM_024643	1.70
spinocerebellar ataxia 8	AF126749	1.70
Consensus includes gb:AL157428.1 /DEF=Homo sapiens mRNA; cDNA DKFZp761F152 (from clone DKFZp761F152). /FEA=mRNA /DB_XREF=gi:7018458 /UG=Hs.306479 Homo sapiens mF	AL157428	1.70
kallikrein 3, (prostate specific antigen)	NM_001648	1.70
Consensus includes gb:BF031829 /FEA=EST /DB_XREF=gi:10739541 /DB_XREF=est:601557983F1 /CLONE=IMAGE:3827709 /UG=Hs.2631 desmoglein 2 /FL=gb:NM_001943.1	BF031829	1.70
hemoglobin, alpha 2	T50399	1.70
catenin (cadherin-associated protein), alpha pseudogene 1	Z37994	1.70
transient receptor potential channel 6	NM_004621	1.70
hypothetical protein DKFZp761D1823	AK026105	1.70

solute carrier family 3 (cystine, dibasic and neutral amino acid transporters, activator of cystine, dibasic and neutral amino acid transport), member 1	NM_000341	1.70
SMART/HDAC1 associated repressor protein	AL524033	1.70
hypothetical protein MGC3020	NM_024048	1.70
erythrocyte membrane protein band 4.1-like 1	AA573523	1.70
neuroligin 3	NM_018977	1.70
Consensus includes gb:N64802 /FEA=EST /DB_XREF=gi:1212631 /DB_XREF=est:yz31b05.s1 /CLONE=IMAGE:284625 /UG=Hs.173684 Homo sapiens mRNA; cDNA DKFZp762G207 (from cl	N64802	1.70
aquaporin 9	NM_020980	1.70
Consensus includes gb:AW450360 /FEA=EST /DB_XREF=gi:6991136 /DB_XREF=est:UI-H-BI3-akn-c-11-0-UI.s1 /CLONE=IMAGE:2734845 /UG=Hs.98834 ESTs	AW450360	1.70
Consensus includes gb:AL117549.1 /DEF=Homo sapiens mRNA; cDNA DKFZp586H0123 (from clone DKFZp586H0123). /FEA=mRNA /DB_XREF=gi:5912083 /UG=Hs.268496 Homo sapiens	AL117549	1.70
FXYD domain-containing ion transport regulator 2	X86400	1.70
gb:U61166.1 /DEF=Human SH3 domain-containing protein SH3P17 mRNA, complete cds. /FEA=mRNA /PROD=SH3 domain-containing protein SH3P17 /DB_XREF=gi:1438932 /UG=Hs.3071	U61166	1.70
melanoma antigen, family A, 6	U10691	1.70
macrophage stimulating 1 receptor (c-met-related tyrosine kinase)	NM_002447	1.70
Consensus includes gb:AI911273 /FEA=EST /DB_XREF=gi:5631009 /DB_XREF=est:wg33f09.x1 /CLONE=IMAGE:2366921 /UG=Hs.122158 ESTs	AI911273	1.70
EGF-like-domain, multiple 5	BF110421	1.70
Consensus includes gb:AL080233.1 /DEF=Homo sapiens mRNA; cDNA DKFZp586L111 (from clone DKFZp586L111). /FEA=mRNA /DB_XREF=gi:5262726 /UG=Hs.326580 Homo sapiens mR	AL080233	1.70
tumor necrosis factor (ligand) superfamily, member 13	AF114012	1.70
cholinephosphotransferase 1	AF195624	1.70
interferon regulatory factor 2	NM_002199	1.70
RNA binding motif, single stranded interacting protein 2	AA536000	1.70
Consensus includes gb:AV700891 /FEA=EST /DB_XREF=gi:10302862 /DB_XREF=est:AV700891 /CLONE=GKCBQD03 /UG=Hs.292477 ESTs	AV700891	1.70
hypothetical protein FLJ20856	NM_025143	1.70
leukocyte immunoglobulin-like receptor, subfamily A (without TM domain), member 4	AF041261	1.70
zinc finger protein 221	NM_013359	1.70
DKFZP564C103 protein	NM_015654	1.69
hypothetical protein FLJ10815	AK022786	1.69
hypothetical protein FLJ12121	NM_024978	1.69
Consensus includes gb:AU147962 /FEA=EST /DB_XREF=gi:11009483 /DB_XREF=est:AU147962 /CLONE=MAMMA1002250 /UG=Hs.301509 Homo sapiens cDNA FLJ12339 fis, clone MAMM	AU147962	1.69
killer cell lectin-like receptor subfamily D, member 1	U30610	1.69
proteoglycan 1, secretory granule	NM_002727	1.69
growth factor independent 1	NM_005263	1.69
Consensus includes gb:AL033538 /DEF=Human DNA sequence from clone RP3-477H23 on chromosome 22q12.1-12.2 Contains parts of one or two novel genes, ESTs, STSs and GSSs /FEA=	AL033538	1.69
calsequestrin 2 (cardiac muscle)	NM_001232	1.69
Consensus includes gb:AL137403.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434L092 (from clone DKFZp434L092). /FEA=mRNA /DB_XREF=gi:6807951 /UG=Hs.306457 Homo sapiens mR	AL137403	1.69
keratocan	NM_007035	1.69
putative peroxisome microbody protein 175.1	NM_015887	1.69
timeless homolog (Drosophila)	AK000721	1.69
ATPase, H+/K+ exchanging, beta polypeptide	NM_000705	1.69
Sjogren syndrome antigen B (autoantigen La)	BG532929	1.69
Consensus includes gb:AL080068.1 /DEF=Homo sapiens mRNA; cDNA DKFZp564J062 (from clone DKFZp564J062). /FEA=mRNA /DB_XREF=gi:5262475 /UG=Hs.272534 Homo sapiens mR	AL080068	1.69
KIAA0984 protein	AW134976	1.69
protein tyrosine phosphatase, receptor type, D	NM_002839	1.69
hypothetical protein FLJ10178	AK001040	1.69
MEF like Protein	NM_020356	1.69
met proto-oncogene (hepatocyte growth factor receptor)	BE870509	1.69
small inducible cytokine A2 (monocyte chemotactic protein 1)	S69738	1.69
transcription elongation factor (SIII) elongin A2	NM_016427	1.69
Consensus includes gb:AF023203.1 /DEF=Homo sapiens homeobox protein Og12 (OGL12) mRNA, partial cds. /FEA=mRNA /GEN=OGL12 /PROD=homeobox protein Og12 /DB_XREF=gi:297	AF023203	1.69
immunoglobulin lambda locus	X93006	1.69
mitochondrial ribosomal protein L44	NM_022915	1.69
v-akt murine thymoma viral oncogene homolog 3 (protein kinase B, gamma)	NM_005465	1.69
KIAA1396 protein	AF352026	1.69
interleukin 18 (interferon-gamma-inducing factor)	NM_001562	1.69
nerve growth factor receptor (TNFRSF16) associated protein 1	NM_014380	1.69
H3 histone family, member D	NM_003532	1.69
DnaJ (Hsp40) homolog, subfamily B, member 6	AL355379	1.69
hypothetical protein FLJ13881	NM_024729	1.69
Consensus includes gb:AK025619.1 /DEF=Homo sapiens cDNA: FLJ21966 fis, clone HEP05644. /FEA=mRNA /DB_XREF=gi:10438192 /UG=Hs.287688 Homo sapiens cDNA: FLJ21966 fis, cl	AK025619	1.69
Consensus includes gb:AL117659.1 /DEF=Homo sapiens mRNA; cDNA DKFZp586I0624 (from clone DKFZp586I0624). /FEA=mRNA /DB_XREF=gi:5912252 /UG=Hs.306353 Homo sapiens n	AL117659	1.69
growth arrest-specific 2	NM_005256	1.69
cartilage acidic protein 1	NM_018058	1.69
mitogen-activated protein kinase 8	U34822	1.69
LAT1-3TM protein	AC003007	1.69

cisplatin resistance-associated overexpressed protein	AW089673	1.69
Consensus includes gb:AK022120.1 /DEF=Homo sapiens cDNA FLJ12058 fis, clone HEMBB1002092. /FEA=mRNA /DB_XREF=gi:10433445 /UG=Hs.296688 Homo sapiens cDNA FLJ12058 f	AK022120	1.69
nuclear receptor subfamily 4, group A, member 2	S77154	1.69
promyelocytic leukemia	AF230410	1.69
thymosin, beta 10	AF090913	1.69
sodium channel, voltage-gated, type XII, alpha polypeptide	AF150882	1.69
malignancy-associated protein	AF041410	1.69
CD33 antigen (gp67)	NM_001772	1.69
Consensus includes gb:AW969675 /FEA=EST /DB_XREF=gi:8159519 /DB_XREF=est:EST381752 /UG=Hs.291232 ESTs	AW969675	1.69
hypothetical protein OVC10-2	NM_018692	1.69
Cdc42 effector protein 3	AI754416	1.69
transport-secretion protein 2.2,	AI979087	1.69
olfactory receptor, family 1, subfamily A, member 1	NM_014565	1.69
Consensus includes gb:AF203728.1 /DEF=Homo sapiens bone morphogenetic protein BMPY (BMPY) mRNA, partial cds. /FEA=mRNA /GEN=BMPY /PROD=bone morphogenetic protein BMP'	AF203728	1.69
laminin, alpha 4	U77706	1.69
leucine rich repeat (in FLII) interacting protein 1	AF130054	1.69
olfactory receptor, family 5, subfamily V member 1	NM_030876	1.69
Consensus includes gb:AW268081 /FEA=EST /DB_XREF=gi:6655111 /DB_XREF=est:xr92e03.x1 /CLONE=IMAGE:2767612 /UG=Hs.193408 ESTs, Weakly similar to ALUC_HUMAN !!!! ALU CL	AW268081	1.69
killer cell immunoglobulin-like receptor, three domains, long cytoplasmic tail, 1	U73396	1.69
Consensus includes gb:AK021474.1 /DEF=Homo sapiens cDNA FLJ11412 fis, clone HEMBA1000876. /FEA=mRNA /DB_XREF=gi:10432666 /UG=Hs.289094 Homo sapiens cDNA FLJ11412 fi	AK021474	1.69
NK3 transcription factor homolog A (Drosophila)	AF247704	1.69
leucine-rich, glioma inactivated 1	NM_005097	1.69
KIAA0346 protein	AA521267	1.69
LOC89231	AK026768	1.69
KIAA1053 protein	AL117523	1.69
gb:BC005043.1 /DEF=Homo sapiens, Similar to tumor necrosis factor receptor superfamily, member 10c, decoy without an intracellular domain, clone MGC:12787, mRNA, complete cds. /FE/	BC005043	1.68
Consensus includes gb:AW663632 /FEA=EST /DB_XREF=gi:7456210 /DB_XREF=est:hj12a03.x1 /CLONE=IMAGE:2981548 /UG=Hs.285625 Homo sapiens mRNA; cDNA DKFZp434A119 (fro	AW663632	1.68
nuclear fragile X mental retardation protein interacting protein 1	AW593143	1.68
KIAA0523 protein	AB011095	1.68
CD36 antigen (collagen type I receptor, thrombospondin receptor)	M98399	1.68
cAMP responsive element binding protein 1	NM_004379	1.68
docking protein 2, 56kD	AI828929	1.68
Kruppel-like factor 4 (gut)	NM_004235	1.68
Consensus includes gb:AV687010 /FEA=EST /DB_XREF=gi:10288873 /DB_XREF=est:AV687010 /CLONE=GKCAYC04 /UG=Hs.284651 ESTs	AV687010	1.68
outer dense fibre of sperm tails 1	NM_024410	1.68
Cdc42 effector protein 4; binder of Rho GTPases 4	AL162074	1.68
Consensus includes gb:AL049263.1 /DEF=Homo sapiens mRNA; cDNA DKFZp564F133 (from clone DKFZp564F133). /FEA=mRNA /DB_XREF=gi:4500011 /UG=Hs.306292 Homo sapiens mF	AL049263	1.68
DKFZP434B204 protein	AL080148	1.68
UV-B repressed sequence, HUR 7	BE148534	1.68
cytochrome P450, subfamily I (aromatic compound-inducible), polypeptide 2	NM_000761	1.68
SRY (sex determining region Y)-box 20	AB025355	1.68
Cluster Incl. AL050166:Homo sapiens mRNA; cDNA DKFZp586D1122 (from clone DKFZp586D1122) /cds=UNKNOWN /gb=AL050166 /gi=4884381 /ug=Hs.26295 /len=2654	AL050166	1.68
crystallin, gamma D	NM_006891	1.68
Consensus includes gb:AW576871 /FEA=EST /DB_XREF=gi:7251920 /DB_XREF=est:RC2-BT0318-110100-012-e02 /UG=Hs.201420 ESTs, Moderately similar to ALU7_HUMAN ALU SUBFAM	AW576871	1.68
KIAA0545 protein	AA425633	1.68
nuclear matrix protein p84	NM_005131	1.68
transducin (beta)-like 1	BF593932	1.68
cytochrome b5 reductase b5R.2	NM_016229	1.68
myeloid/lymphoid or mixed-lineage leukemia (trithorax homolog, Drosophila); translocated to, 2	AA715041	1.68
glutamate-ammonia ligase (glutamine synthase)	U08626	1.68
gb:L14458.1 /DEF=Human Ig rearranged kappa-chain gene V-J-region, complete cds. /FEA=mRNA /DB_XREF=gi:398492 /FL=gb:L14458.1	L14458	1.68
hypothetical protein FLJ21302	NM_022901	1.68
Consensus includes gb:U95737 /DEF=Human Chromosome 16 BAC clone CIT987SK-A-388D4 /FEA=mRNA /DB_XREF=gi:3417290 /UG=Hs.177961 Human Chromosome 16 BAC clone CIT987	U95737	1.68
chemokine (C-C motif) receptor 3	NM_001837	1.68
DKFZp434P211 protein	NM_014549	1.68
hypothetical protein FLJ13110	BE535746	1.68
drebrin 1	AL110225	1.68
killer cell immunoglobulin-like receptor, three domains, long cytoplasmic tail, 2	L76666	1.68
growth differentiation factor 5 (cartilage-derived morphogenetic protein-1)	NM_000557	1.68
hepatocyte nuclear factor 4, gamma	NM_004133	1.68
immunoglobulin heavy constant gamma 3 (G3m marker)	AI275469	1.68
potassium voltage-gated channel, KQT-like subfamily, member 4	NM_004700	1.68
heat shock 70kD protein 2	U56725	1.68

prostaglandin F receptor (FP)	NM_000959	1.68
H1 histone family, member 4	NM_005321	1.68
neighbor of A-kinase anchoring protein 95	AB015332	1.68
solute carrier family 2 (facilitated glucose transporter), member 10	NM_030777	1.68
arginine vasopressin receptor 1B	NM_000707	1.68
synaptopodin	NM_007286	1.68
forkhead box E1 (thyroid transcription factor 2)	NM_004473	1.68
protein tyrosine phosphatase, non-receptor type substrate 1	AC004832	1.68
gelsolin (amyloidosis, Finnish type)	BE675337	1.68
spindlin-like	AF356353	1.68
Consensus includes gb:AK024568.1 /DEF=Homo sapiens cDNA: FLJ20915 fis, clone ADSE00692. /FEA=mRNA /DB_XREF=gi:10436878 /UG=Hs.306691 Homo sapiens cDNA: FLJ20915 fis, c	AK024568	1.68
Consensus includes gb:AF070579.1 /DEF=Homo sapiens clone 24487 mRNA sequence. /FEA=mRNA /DB_XREF=gi:3387951 /UG=Hs.283819 Homo sapiens clone 24487 mRNA sequence	AF070579	1.68
surfactant, pulmonary-associated protein D	NM_003019	1.68
kinesin family member 5A	NM_004984	1.68
hypothetical protein FLJ11016	NM_018301	1.68
hypothetical protein FLJ10891	NM_018260	1.68
ATPase, aminophospholipid transporter-like, Class I, type 8A, member 2	NM_016529	1.67
statherin	NM_003154	1.67
Consensus includes gb:BE888744 /FEA=EST /DB_XREF=gi:10345354 /DB_XREF=est:601513656F1 /CLONE=IMAGE:3915133 /UG=Hs.169274 ESTs, Highly similar to IFT2_HUMAN INTERFI	BE888744	1.67
cytochrome P450, subfamily IIA (phenobarbital-inducible), polypeptide 6	T67741	1.67
Consensus includes gb:AK021930.1 /DEF=Homo sapiens cDNA FLJ11868 fis, clone HEMBA1006993. /FEA=mRNA /DB_XREF=gi:10433231 /UG=Hs.302260 Homo sapiens cDNA FLJ11868 f	AK021930	1.67
platelet-derived growth factor receptor, alpha polypeptide	M22734	1.67
chromosome 4 open reading frame 6	NM_005750	1.67
TYRO3 protein tyrosine kinase	D50479	1.67
tropomyosin 1 (alpha)	AA923707	1.67
hypothetical protein FLJ14001	NM_024677	1.67
lysyl oxidase	BE503425	1.67
CD1B antigen, b polypeptide	NM_001764	1.67
tryptase beta 1	NM_003294	1.67
activating transcription factor 7	NM_006856	1.67
SMAD in the antisense orientation	NM_022001	1.67
a disintegrin and metalloproteinase domain 5	AI132823	1.67
fms-related tyrosine kinase 3	NM_004119	1.67
hyaluronoglucosaminidase 4	NM_012269	1.67
hypothetical protein FLJ12800	NM_022903	1.67
immunoglobulin lambda-like polypeptide 2	L02326	1.67
recombining binding protein suppressor of hairless (Drosophila)-like	NM_014276	1.67
Consensus includes gb:AK021988.1 /DEF=Homo sapiens cDNA FLJ11926 fis, clone HEMBB1000374. /FEA=mRNA /DB_XREF=gi:10433297 /UG=Hs.306617 Homo sapiens cDNA FLJ11926 f	AK021988	1.67
exostoses (multiple)-like 1	NM_004455	1.67
cytochrome P450, subfamily VIIB (oxysterol 7 alpha-hydroxylase), polypeptide 1	NM_004820	1.67
melanoma antigen, family C, 1	NM_005462	1.67
SEC14-like 2 (S. cerevisiae)	NM_012429	1.67
trophoblast glycoprotein	NM_006670	1.67
Consensus includes gb:AK024525.1 /DEF=Homo sapiens cDNA: FLJ20872 fis, clone ADKA02604. /FEA=mRNA /DB_XREF=gi:10436827 /UG=Hs.306683 Homo sapiens cDNA: FLJ20872 fis, c	AK024525	1.67
Consensus includes gb:AU148086 /FEA=EST /DB_XREF=gi:11009607 /DB_XREF=est:AU148086 /CLONE=MAMMA1002554 /UG=Hs.302691 Homo sapiens cDNA FLJ12379 fis, clone MAMM	AU148086	1.67
DKFZP586J0119 protein	AF112207	1.67
tumor necrosis factor receptor superfamily, member 11a, activator of NFKB	NM_003839	1.67
gonadotropin-releasing hormone 2	NM_001501	1.67
HSPC073 protein	NM_014163	1.67
hypothetical protein FLJ22215	NM_022834	1.67
cytochrome P450, subfamily IIIA (niphedipine oxidase), polypeptide 3	NM_000776	1.67
Consensus includes gb:AK026820.1 /DEF=Homo sapiens cDNA: FLJ23167 fis, clone LNG09902. /FEA=mRNA /DB_XREF=gi:10439765 /UG=Hs.287730 Homo sapiens cDNA: FLJ23167 fis, cl	AK026820	1.67
epidermal growth factor (beta-urogastrone)	NM_001963	1.67
nuclear respiratory factor 1	AW003022	1.67
Consensus includes gb:AA634446 /FEA=EST /DB_XREF=gi:2557660 /DB_XREF=est:zu79e01.s1 /CLONE=IMAGE:744216 /UG=Hs.116840 ESTs	AA634446	1.67
LIM domain binding 2	NM_001290	1.67
ral guanine nucleotide dissociation stimulator	AI421559	1.67
likely ortholog of mouse gangfioside-induced differentiation-associated-protein 2	NM_017686	1.67
UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase 3 (GalNAc-T3)	NM_004482	1.67
Consensus includes gb:AL049285.1 /DEF=Homo sapiens mRNA; cDNA DKFZp564M193 (from clone DKFZp564M193). /FEA=mRNA /DB_XREF=gi:4500045 /UG=Hs.302053 Homo sapiens m	AL049285	1.67
Consensus includes gb:AA479678 /FEA=EST /DB_XREF=gi:2205564 /DB_XREF=est:zu43a04.s1 /CLONE=IMAGE:740718 /UG=Hs.203269 ESTs, Moderately similar to ALU8_HUMAN ALU SU	AA479678	1.67
jumonji homolog (mouse)	BG029530	1.67
ATPase, Cu++ transporting, alpha polypeptide (Menkes syndrome)	NM_000052	1.67

melanoma antigen, family C, 1	AL022152	1.67
laminin, beta 1	NM_002291	1.67
neuropeptide Y	NM_000905	1.67
tumor necrosis factor (ligand) superfamily, member 10	U57059	1.67
Consensus includes gb:BE046461 /FEA=EST /DB_XREF=gi:8363514 /DB_XREF=est:hn47g05.x2 /CLONE=IMAGE:3026840 /UG=Hs.159412 Homo sapiens clone 24628 mRNA sequence	BE046461	1.67
E3 ubiquitin ligase SMURF2	AY014180	1.67
osteomodulin	NM_005014	1.67
occludin	U53823	1.67
plakophilin 3	BF001503	1.67
Consensus includes gb:AL050042.1 /DEF=Homo sapiens mRNA; cDNA DKFZp566L0824 (from clone DKFZp566L0824). /FEA=mRNA /DB_XREF=gi:4884284 /UG=Hs.306309 Homo sapiens	AL050042	1.67
Consensus includes gb:AW474615 /FEA=EST /DB_XREF=gi:7044721 /DB_XREF=est:xy18c02.x1 /CLONE=IMAGE:2853506 /UG=Hs.250988 ESTs	AW474615	1.67
phospholysine phosphohistidine inorganic pyrophosphate phosphatase	NM_022126	1.67
discoidin domain receptor family, member 2	NM_006182	1.67
forkhead box C2 (MFH-1, mesenchyme forkhead 1)	NM_005251	1.67
gamma-glutamyltransferase 1	J05235	1.67
hypothetical protein PRO1094	NM_018575	1.67
phosphoinositide-3-kinase, catalytic, beta polypeptide	AA805318	1.67
cytochrome P450, subfamily IVF, polypeptide 2	D26480	1.67
Consensus includes gb:AW085172 /FEA=EST /DB_XREF=gi:6040324 /DB_XREF=est:xe06g09.x1 /CLONE=IMAGE:2606368 /UG=Hs.243366 ESTs, Highly similar to KPCM_HUMAN PROTEIN I	AW085172	1.67
secreted frizzled-related protein 4	AW089415	1.67
Consensus includes gb:AF308291.1 /DEF=Homo sapiens serologically defined breast cancer antigen NY-BR-46 mRNA, partial cds. /FEA=mRNA /PROD=serologically defined breast cancer a	AF308291	1.67
aggrecan 1 (chondroitin sulfate proteoglycan 1, large aggregating proteoglycan, antigen identified by monoclonal antibody A0122)	NM_013227	1.67
Consensus includes gb:AA995925 /FEA=EST /DB_XREF=gi:3182414 /DB_XREF=est:ou93g10.s1 /CLONE=IMAGE:1635426 /UG=Hs.7309 Homo sapiens clone 23741 mRNA sequence	AA995925	1.67
hypothetical protein	AJ740515	1.67
phosphodiesterase 11A	NM_016953	1.67
Consensus includes gb:AL080112.1 /DEF=Homo sapiens mRNA; cDNA DKFZp586H0722 (from clone DKFZp586H0722). /FEA=mRNA /DB_XREF=gi:5262539 /UG=Hs.332731 Homo sapiens	AL080112	1.67
TNF receptor-associated factor 3	NM_003300	1.67
hypothetical protein FLJ21240	NM_024847	1.67
caspase recruitment domain protein 10	AW205153	1.67
POU domain, class 4, transcription factor 3	NM_002700	1.66
cyclin-dependent kinase inhibitor 1C (p57, Kip2)	N95363	1.66
Consensus includes gb:AA100250 /FEA=EST /DB_XREF=gi:1647823 /DB_XREF=est:zn41d10.s1 /CLONE=IMAGE:550003 /UG=Hs.90869 Homo sapiens clones 24622 and 24623 mRNA seq	AA100250	1.66
fascin (Strongylocentrotus purpuratus) homolog 3 (actin-bundling protein, testicular)	NM_020369	1.66
enolase 2, (gamma, neuronal)	NM_001975	1.66
hairless homolog (mouse)	AF039196	1.66
p53-regulated apoptosis-inducing protein 1	NM_022112	1.66
hypothetical protein FLJ10633	NM_018162	1.66
KIAA0368 protein	AI272899	1.66
gastrin-releasing peptide	NM_002091	1.66
tumor necrosis factor receptor superfamily, member 14 (herpesvirus entry mediator)	BC002794	1.66
U6 snRNA-associated Sm-like protein	N37057	1.66
zinc finger protein 258	NM_007167	1.66
hemochromatosis	AF144240	1.66
SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily e, member 1	BG289800	1.66
spermidine/spermine N1-acetyltransferase	M55580	1.66
wingless-type MMTV integration site family, member 2B	AB045117	1.66
lipase protein	AA209239	1.66
Sam68-like phosphotyrosine protein, T-STAR	AF069681	1.66
hypothetical protein	NM_018992	1.66
KIAA0455 gene product	AW592563	1.66
KIAA0669 gene product	AF201291	1.66
class I cytokine receptor	AI983115	1.66
weakly similar to glutathione peroxidase 2	AA406605	1.66
SMT3 suppressor of mif two 3 homolog 2 (yeast)	AV726646	1.66
mannan-binding lectin serine protease 1 (C4/C2 activating component of Ra-reactive factor)	AV686235	1.66
calcium channel, voltage-dependent, L type, alpha 1D subunit	BE550599	1.66
myelodysplasia syndrome 1	NM_004991	1.66
neuronal specific transcription factor DAT1	AL050152	1.66
Consensus includes gb:BF155672 /FEA=EST /DB_XREF=gi:11050855 /DB_XREF=est:QV4-HT0894-290900-399-c04 /UG=Hs.146200 ESTs	BF155672	1.66
ubiquitin associated and SH3 domain containing, A	NM_018961	1.66
Consensus includes gb:AL031118 /DEF=Human DNA sequence from clone 153G14 on chromosome 6p21.3-22.2. Contains the 3 end of the ZNF184 gene for Kruppel-like zinc finger protein	AL031118	1.66
zinc finger protein 37 homolog (mouse)	NM_003408	1.66
Consensus includes gb:AI344415 /FEA=EST /DB_XREF=gi:4081621 /DB_XREF=est:qp13b08.x1 /CLONE=IMAGE:1917879 /UG=Hs.156082 ESTs	AI344415	1.66

PTEN induced putative kinase 1	AF316873	1.66
keratin 7	AI920979	1.66
hypothetical protein DKFZp761N09121	BF435376	1.66
capicua homolog (Drosophila)	AV755522	1.66
Consensus includes gb:BG109746 /FEA=EST /DB_XREF=gi:12603252 /DB_XREF=est:602280883F1 /CLONE=IMAGE:4368381 /UG=Hs.325625 Homo sapiens clone 23938 mRNA sequence	BG109746	1.66
Consensus includes gb:AF090910.1 /DEF=Homo sapiens clone HQ0262. /FEA=mRNA /DB_XREF=gi:6690195 /UG=Hs.236760 Homo sapiens clone HQ0262	AF090910	1.66
integrin, beta 1 (fibronectin receptor, beta polypeptide, antigen CD29 includes MDF2, MSK12)	M34189	1.66
glutamate receptor, metabotropic 3	NM_000840	1.66
hypothetical protein FLJ21665	NM_024803	1.66
hypothetical protein FLJ20294	AB051523	1.66
2'-5'-oligoadenylate synthetase-like	AF063612	1.66
integrin, beta 5	AI335208	1.66
deleted in lymphocytic leukemia, 2	AF264787	1.66
prolactin receptor	AF166329	1.66
tyrosine kinase with immunoglobulin and epidermal growth factor homology domains	NM_005424	1.66
tumor protein D52-like 1	NM_003287	1.66
ameloblastin, enamel matrix protein	NM_016519	1.66
myeloid/lymphoid or mixed-lineage leukemia (trithorax homolog, Drosophila); translocated to, 4	AL049698	1.66
zinc finger protein-like 1	AI401287	1.66
atonal homolog 1 (Drosophila)	NM_005172	1.66
glucokinase (hexokinase 4, maturity onset diabetes of the young 2)	M69051	1.66
forkhead box O1A (rhabdomyosarcoma)	AW117498	1.66
sucrase-isomaltase	NM_001041	1.66
small proline-rich protein 1A	NM_005987	1.66
Consensus includes gb:AF113683.1 /DEF=Homo sapiens clone FLB3521. /FEA=mRNA /DB_XREF=gi:6855610 /UG=Hs.306569 Homo sapiens clone FLB3521	AF113683	1.66
KIAA0285 gene product	NM_014807	1.66
aristaless homeobox (Drosophila)	AI469991	1.66
KIAA0863 protein	AK022688	1.66
hypothetical protein FLJ12668	NM_024997	1.66
hypothetical protein FLJ11807	NM_024954	1.66
RalP2 interacting protein B	NM_006695	1.66
hypothetical protein FLJ12616	NM_024995	1.66
transition protein 1 (during histone to protamine replacement)	NM_003284	1.66
hypothetical protein FLJ11952	AK022014	1.66
connexin46.6	NM_020435	1.66
wingless-type MMTV integration site family, member 3	NM_030753	1.66
potassium inwardly-rectifying channel, subfamily J, member 3	NM_002239	1.66
period homolog 1 (Drosophila)	NM_002616	1.66
gb:BC006381.1 /DEF=Homo sapiens, clone MGC:13287, mRNA, complete cds. /FEA=mRNA /PROD=Unknown (protein for MGC:13287) /DB_XREF=gi:13623544 /FL=gb:BC006381.1	BC006381	1.66
nucleophosmin (nucleolar phosphoprotein B23, numatrin)	AL353580	1.66
dual oxidase 2	NM_014080	1.66
similar to sialyltransferase 7 ((alpha-N-acetylneuraminyl 2,3-betagalactosyl-1,3)-N-acetyl galactosaminide alpha-2,6-sialyltransferase) E	NM_030965	1.65
hypothetical protein FLJ20619	NM_017904	1.65
caveolin 2	BF197655	1.65
Consensus includes gb:AV699347 /FEA=EST /DB_XREF=gi:10301318 /DB_XREF=est:AV699347 /CLONE=GKCBPH07 /UG=Hs.83623 nuclear receptor subfamily 1, group I, member 3	AV699347	1.65
Consensus includes gb:AI979276 /FEA=EST /DB_XREF=gi:5804295 /DB_XREF=est:wu03a04.x1 /CLONE=IMAGE:2515854 /UG=Hs.177482 ESTs, Weakly similar to KIAA1276 protein H.sapie	AI979276	1.65
glutaminase	AF097493	1.65
killer cell immunoglobulin-like receptor, two domains, long cytoplasmic tail, 4	AF002256	1.65
Consensus includes gb:AL050002.1 /DEF=Homo sapiens mRNA; cDNA DKFZp564O222 (from clone DKFZp564O222). /FEA=mRNA /DB_XREF=gi:4884256 /UG=Hs.94795 Homo sapiens mRI	AL050002	1.65
Consensus includes gb:AW405975 /FEA=EST /DB_XREF=gi:6925032 /DB_XREF=est:UI-HF-BL0-acv-a-05-0-UI.r1 /CLONE=IMAGE:3060368 /UG=Hs.247721 Homo sapiens clone mcg53-54 i	AW405975	1.65
hypothetical protein FLJ20519	NM_017860	1.65
calsyntenin-2	NM_022131	1.65
variable charge protein on X with two repeats	NM_016378	1.65
Fc fragment of IgA, receptor for	U56236	1.65
KIAA0632 protein	NM_015545	1.65
hypothetical protein FLJ11112	NM_018327	1.65
Consensus includes gb:AA126763 /FEA=EST /DB_XREF=gi:1686263 /DB_XREF=est:zn87b11.s1 /CLONE=IMAGE:565149 /UG=Hs.182296 ESTs, Weakly similar to ALU6_HUMAN ALU SUBFAI	AA126763	1.65
transcription elongation factor A (SII)-like 1	NM_004780	1.65
hypothetical protein FLJ20307	NM_017757	1.65
bassoon (presynaptic cytomatrix protein)	NM_003458	1.65
Ria	NM_016563	1.65
carboxypeptidase A1 (pancreatic)	NM_001868	1.65
CD163 antigen	Z22969	1.65

mitogen-activated protein kinase 8 interacting protein 3	AB028989	1.65
tripartite motif-containing 10	AF220123	1.65
paired mesoderm homeo box 1	M95929	1.65
glutamate dehydrogenase 1	NM_005271	1.65
Pirin	NM_003662	1.65
kinesin family member 1C	AB014606	1.65
lectin, galactoside-binding, soluble, 4 (galectin 4)	NM_006149	1.65
hypothetical protein MGC10771	NM_024506	1.65
myosin IB	BF432550	1.65
secreted frizzled-related protein 4	NM_003014	1.65
special AT-rich sequence binding protein 1 (binds to nuclear matrix/scaffold-associating DNA's)	NM_002971	1.65
annexin A6	NM_001155	1.65
potassium inwardly-rectifying channel, subfamily J, member 15	U73191	1.65
tumor protein D52-like 1	AF208012	1.65
tight junction protein 1 (zona occludens 1)	AA813018	1.65
serine racemase	NM_021947	1.65
nerve growth factor, beta polypeptide	NM_002506	1.65
ephrin-A3	AW189015	1.65
retinitis pigmentosa GTPase regulator	NM_000328	1.65
zinc finger protein 358	NM_018083	1.65
v-myc myelocytomatosis viral related oncogene, neuroblastoma derived (avian)	AF320053	1.65
histamine receptor H1	NM_000861	1.65
chromosome 5 open reading frame 4	H93077	1.65
S88I31 protein	NM_014035	1.65
SRY (sex determining region Y)-box 5	NM_006940	1.65
gamma-aminobutyric acid (GABA) B receptor, 1	AL031983	1.65
vinculin	NM_014000	1.65
Consensus includes gb:AK024995.1 /DEF=Homo sapiens cDNA: FLJ21342 fis, clone COL02673. /FEA=mRNA /DB_XREF=gi:10437426 /UG=Hs.306739 Homo sapiens cDNA: FLJ21342 fis, cl	AK024995	1.65
nestin (Reed-Steinberg cell-expressed intermediate filament-associated protein)	NM_002956	1.65
hypothetical protein FLJ20712	NM_017937	1.65
cylindromatosis (turban tumor syndrome)	AJ453099	1.65
Consensus includes gb:AL110206.1 /DEF=Homo sapiens mRNA; cDNA DKFZp586N2022 (from clone DKFZp586N2022). /FEA=mRNA /DB_XREF=gi:5817125 /UG=Hs.306339 Homo sapiens	AL110206	1.65
Consensus includes gb:AL117571.1 /DEF=Homo sapiens mRNA; cDNA DKFZp564O1172 (from clone DKFZp564O1172). /FEA=mRNA /DB_XREF=gi:5912123 /UG=Hs.241562 Homo sapiens	AL117571	1.65
Consensus includes gb:AK024784.1 /DEF=Homo sapiens cDNA: FLJ21131 fis, clone CAS06355. /FEA=mRNA /DB_XREF=gi:10437168 /UG=Hs.28264 Homo sapiens mRNA; cDNA DKFZp564I	AK024784	1.65
DKFZP564I922 protein	AF245505	1.65
testicular haploid expressed gene	NM_016585	1.65
cholinergic receptor, nicotinic, beta polypeptide 4	NM_000750	1.65
protein tyrosine phosphatase, receptor type, N polypeptide 2	NM_002847	1.65
tight junction protein 3 (zona occludens 3)	NM_014428	1.65
transforming growth factor, beta-induced, 68kD	NM_000358	1.65
potassium voltage-gated channel, shaker-related subfamily, member 5	NM_002234	1.65
melanoma antigen recognized by T cells 2	NM_018194	1.65
hypothetical protein FLJ11585	BF476502	1.65
deleted in colorectal carcinoma	NM_005215	1.65
oxysterol-binding protein-related protein 1	NM_018030	1.65
p21(CDKN1A)-activated kinase 4	AB032968	1.65
Consensus includes gb:AL110223.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434M063 (from clone DKFZp434M063). /FEA=mRNA /DB_XREF=gi:5817158 /UG=Hs.274521 Homo sapiens m	AL110223	1.65
inhibitor of growth family, member 1-like	NM_001564	1.65
myeloid/lymphoid or mixed-lineage leukemia (trithorax homolog, Drosophila); translocated to, 4	AB011399	1.65
Consensus includes gb:AF023203.1 /DEF=Homo sapiens homeobox protein Og12 (OGL12) mRNA, partial cds. /FEA=mRNA /GEN=OGL12 /PROD=homeobox protein Og12 /DB_XREF=gi:297	AF023203	1.65
paraoxonase 2	NM_000305	1.65
Consensus includes gb:AW149827 /FEA=EST /DB_XREF=gi:6197723 /DB_XREF=est:xf42g03.x1 /CLONE=IMAGE:2620756 /UG=Hs.293272 ESTs, Weakly similar to ALUF_HUMAN !!!! ALU CL	AW149827	1.65
hypothetical protein FLJ12619	BG252842	1.65
transmembrane phosphatase with tensin homology	NM_013315	1.65
LBP protein; likely ortholog of mouse CRTR-1	NM_014553	1.65
fibroblast growth factor receptor 2 (bacteria-expressed kinase, keratinocyte growth factor receptor, craniofacial dysostosis 1, Crouzon syndrome, Pfeiffer syndrome, Jackson-Weiss syndrome	M80634	1.65
SH3-domain, GRB2-like, endophilin B2	BC005407	1.65
runt-related transcription factor 2	L40992	1.65
ectodermal dysplasia 1, anhidrotic	AF061193	1.65
protein phosphatase 1, regulatory (inhibitor) subunit 3A (glycogen and sarcoplasmic reticulum binding subunit, skeletal muscle)	NM_002711	1.65
aldehyde dehydrogenase 3 family, member A2	NM_000382	1.65
type 1 protein phosphatase inhibitor	NM_025210	1.65
TYRO3 protein tyrosine kinase	U05682	1.65

thymosin, beta, identified in neuroblastoma cells	NM_021992	1.65
hypothetical protein PRO2521	NM_018530	1.65
solute carrier family 12 (potassium/chloride transporters), member 4	AI817690	1.65
novel putative protein similar to YIL091C yeast hypothetical 84 kD protein from SGA1-KTR7	NM_014388	1.65
hemopoietic cell kinase	NM_002110	1.65
PDZ-73 protein	AB006955	1.64
caldesmon 1	NM_004342	1.64
tubulin, beta, 5	AL567012	1.64
endothelial PAS domain protein 1	NM_001430	1.64
hypothetical protein FLJ10675	NM_018176	1.64
KIAA0161 gene product	NM_014746	1.64
B melanoma antigen	NM_001187	1.64
KIAA0918 protein	AW449813	1.64
solute carrier family 1 (high affinity aspartate/glutamate transporter), member 6	NM_005071	1.64
apolipoprotein L, 2	NM_030882	1.64
Consensus includes gb:M96936 /DEF=Homo sapiens cystic fibrosis transmembrane conductance regulator (CFTR) gene, exons 23, 24a, and 24 /FEA=mRNA /DB_XREF=gi:180293 /UG=Hs.	M96936	1.64
Consensus includes gb:AA493300 /FEA=EST /DB_XREF=gi:2223141 /DB_XREF=est:ng97e11.s1 /CLONE=IMAGE:942764 /UG=Hs.104258 Homo sapiens mRNA; exon 1, 2, 3, 4, clone:RES4	AA493300	1.64
neurotrophic tyrosine kinase, receptor, type 3	S76475	1.64
A kinase (PRKA) anchor protein (gravin) 12	AB003476	1.64
v-yes-1 Yamaguchi sarcoma viral oncogene homolog 1	NM_005433	1.64
HLA-B associated transcript 1	AI282485	1.64
uronyl 2-sulfotransferase	AW418882	1.64
STAT induced STAT inhibitor 3	AI922872	1.64
lethal (3) malignant brain tumor l(3)mbt protein (Drosophila) homolog	NM_015478	1.64
Consensus includes gb:BF511164 /FEA=EST /DB_XREF=gi:11594462 /DB_XREF=est:UI-H-BI4-aoi-a-09-0-UI.s1 /CLONE=IMAGE:3084952 /UG=Hs.293309 ESTs	BF511164	1.64
Rho GTPase activating protein 5	AW044631	1.64
arachidonate 12-lipoxygenase, 12R type	NM_001139	1.64
hypothetical protein LOC57158	NM_020433	1.64
Nck, Ash and phospholipase C binding protein	AI968839	1.64
Consensus includes gb:AI351043 /FEA=EST /DB_XREF=gi:4088249 /DB_XREF=est:qt22a08.x1 /CLONE=IMAGE:1948310 /UG=Hs.279477 ESTs	AI351043	1.64
putative X-linked retinopathy protein	S58722	1.64
protocadherin gamma subfamily C, 3	AF152524	1.64
EphB1	AF037334	1.64
GTP binding protein overexpressed in skeletal muscle	NM_005261	1.64
potassium voltage-gated channel, Shaw-related subfamily, member 4	NM_004978	1.64
adrenomedullin receptor	NM_007264	1.64
Consensus includes gb:BF511231 /FEA=EST /DB_XREF=gi:11594529 /DB_XREF=est:UI-H-BI4-aoi-g-11-0-UI.s1 /CLONE=IMAGE:3085244 /UG=Hs.288582 ESTs, Weakly similar to ubiquitou	BF511231	1.64
caspase 1, apoptosis-related cysteine protease (interleukin 1, beta, convertase)	AI719655	1.64
serine (or cysteine) proteinase inhibitor, clade B (ovalbumin), member 9	BC002538	1.64
SEC22 vesicle trafficking protein-like 1 (S. cerevisiae)	BC001364	1.64
hypothetical protein FLJ22688	NM_025129	1.64
hypothetical protein FLJ23412	NM_022082	1.64
Consensus includes gb:BF447105 /FEA=EST /DB_XREF=gi:11512243 /DB_XREF=est:7p49d07.x1 /CLONE=IMAGE:3649189 /UG=Hs.281706 sortilin 1	BF447105	1.64
H4 histone family, member C	NM_003540	1.64
neuronal PAS domain protein 2	U77970	1.64
CSR1 protein	NM_016240	1.64
a disintegrin and metalloproteinase domain 3a (cyritestin 1)	X89655	1.64
beta-amyloid binding protein precursor	AA012917	1.64
G protein-coupled receptor 31	NM_005299	1.64
voltage-gated sodium channel beta-3 subunit (scn3b gene)	AB032984	1.64
cyclin-E binding protein 1	NM_016323	1.64
hypothetical protein FLJ12650	NM_024522	1.64
hypothetical protein MGC4093	NM_030578	1.64
solute carrier family 21 (organic anion transporter), member 9	AF205073	1.64
zinc finger protein 291	AK025663	1.64
desmoplakin (DPI, DPII)	BE546897	1.64
Consensus includes gb:U00956.1 /DEF=Human clone KDB5.1 (CAC)n(GTG)n repeat-containing mRNA. /FEA=mRNA /DB_XREF=gi:405058 /UG=Hs.104112 Human clone KDB5.1 (CAC)n(GTG	U00956	1.64
hypothetical gene CG018	AL049786	1.64
A kinase (PRKA) anchor protein 10	NM_007202	1.64
cancer/testis antigen	AF038567	1.64
KDEL (Lys-Asp-Glu-Leu) endoplasmic reticulum protein retention receptor 3	NM_006855	1.64
Consensus includes gb:U86813.1 /DEF=Homo sapiens serotonin-7 receptor pseudogene, complete sequence. /FEA=mRNA /PROD=serotonin-7 receptor /DB_XREF=gi:3138916 /UG=Hs.23	U86813	1.64
Consensus includes gb:AK001127.1 /DEF=Homo sapiens cDNA FLJ10265 fis, clone HEMBB1001014. /FEA=mRNA /DB_XREF=gi:7022192 /UG=Hs.255369 Homo sapiens cDNA FLJ10265 fis	AK001127	1.64

Consensus includes gb:AL157837 /DEF=Human DNA sequence from clone RP11-467I21 on chromosome 1q31.1-31.3 Contains a ribosomal protein L23A pseudogene, ESTs, STSs and GSSs /	AL157837	1.64
Consensus includes gb:AI206888 /FEA=EST /DB_XREF=gi:3765560 /DB_XREF=est:qr35a04.x1 /CLONE=IMAGE:1942830 /UG=Hs.154131 ESTs	AI206888	1.64
hypothetical protein FLJ10375	NM_018075	1.64
Consensus includes gb:AL133618.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434C2021 (from clone DKFZp434C2021). /FEA=mRNA /DB_XREF=gi:6599233 /UG=Hs.306443 Homo sapiens	AL133618	1.64
tachykinin, precursor 1 (substance K, substance P, neurokinin 1, neurokinin 2, neuromedin L, neurokinin alpha, neuropeptide K,neuropeptide gamma)	NM_003182	1.64
D-aspartate oxidase	NM_003649	1.64
UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase 8 (GalNAc-T8)	NM_017417	1.64
leukocyte immunoglobulin-like receptor, subfamily B (with TM and ITIM domains), member 3	AF009635	1.64
deleted in liver cancer 1	AF026219	1.64
ACO for serine protease homologue	NM_017509	1.64
dopamine receptor D1	X58987	1.64
solute carrier family 12, (potassium/chloride transporter) member 5	AF208159	1.64
KIAA0090 protein	D42044	1.64
hypothetical protein FLJ12294	NM_025100	1.64
chromobox homolog 5 (HP1 alpha homolog, Drosophila)	L07515	1.64
Consensus includes gb:AL050370.1 /DEF=Homo sapiens mRNA; cDNA DKFZp566C0546 (from clone DKFZp566C0546). /FEA=mRNA /DB_XREF=gi:4914605 /UG=Hs.59728 Homo sapiens m	AL050370	1.64
plasminogen-like	BC005379	1.64
CCR4-NOT transcription complex, subunit 2	BG105204	1.64
caspase recruitment domain family, member 9	NM_022352	1.64
Consensus includes gb:AK026439.1 /DEF=Homo sapiens cDNA: FLJ22786 fis, clone KAIA2150. /FEA=mRNA /DB_XREF=gi:10439304 /UG=Hs.306848 Homo sapiens cDNA: FLJ22786 fis, clo	AK026439	1.64
hypothetical protein FLJ12691	NM_024572	1.64
hyaluronan synthase 1	NM_001523	1.64
transducer of ERBB2, 2	AB051450	1.64
pancreas-enriched phospholipase C	NM_016341	1.64
fibroblast growth factor receptor 2 (bacteria-expressed kinase, keratinocyte growth factor receptor, craniofacial dysostosis 1, Crouzon syndrome, Pfeiffer syndrome, Jackson-Weiss syndrome)	NM_022971	1.64
kinase suppressor of ras	U43586	1.64
chromosome 17 open reading frame 1A	NM_006382	1.64
Consensus includes gb:AF086790.1 /DEF=Homo sapiens aconitase precursor (ACON) mRNA, nuclear gene encoding mitochondrial protein, partial cds. /FEA=mRNA /GEN=ACON /PROD=aco	AF086790	1.64
VPS10 domain receptor protein	AB028982	1.64
Consensus includes gb:AA653456 /FEA=EST /DB_XREF=gi:2589627 /DB_XREF=est:ag66c08.s1 /CLONE=IMAGE:1127918 /UG=Hs.270495 ESTs, Weakly similar to ubiquitous TPR motif, Y i	AA653456	1.64
discs, large (Drosophila) homolog 4	NM_001365	1.64
Consensus includes gb:AK024995.1 /DEF=Homo sapiens cDNA: FLJ21342 fis, clone COL02673. /FEA=mRNA /DB_XREF=gi:10437426 /UG=Hs.306739 Homo sapiens cDNA: FLJ21342 fis, cl	AK024995	1.64
erythrocyte membrane protein band 4.2	M30646	1.64
clg01 protein	NM_014477	1.64
Consensus includes gb:AL110201.1 /DEF=Homo sapiens mRNA; cDNA DKFZp586F1622 (from clone DKFZp586F1622). /FEA=mRNA /DB_XREF=gi:5817120 /UG=Hs.278435 Homo sapiens	AL110201	1.64
microtubule-associated protein 1A	AW296788	1.64
hypothetical protein FLJ12586	NM_024620	1.64
Consensus includes gb:AI697803 /FEA=EST /DB_XREF=gi:4985703 /DB_XREF=est:we17e05.x1 /CLONE=IMAGE:2341376 /UG=Hs.114963 Homo Sapiens (clone B383E13) chromosome 4p	AI697803	1.64
guanine nucleotide binding protein (G protein), q polypeptide	BF222895	1.63
a disintegrin and metalloproteinase domain 20	NM_003814	1.63
KIAA1096 protein	AW301305	1.63
Consensus includes gb:AK026720.1 /DEF=Homo sapiens cDNA: FLJ23067 fis, clone LNG04993. /FEA=mRNA /DB_XREF=gi:10439638 /UG=Hs.117167 Homo sapiens cDNA: FLJ23067 fis, cl	AK026720	1.63
MRE11 meiotic recombination 11 homolog A (S. cerevisiae)	BC005241	1.63
Consensus includes gb:X81637.1 /DEF=H.sapiens clathrin light chain b gene. /FEA=mRNA /DB_XREF=gi:963046 /UG=Hs.73919 clathrin, light polypeptide (Lcb)	X81637	1.63
glycophorin A (includes MN blood group)	U00178	1.63
g20 protein	NM_016210	1.63
desmocollin 3	AI797281	1.63
secretogranin III	NM_013243	1.63
B-cell CLL/lymphoma 3	NM_005178	1.63
hypothetical protein DKFZp564D0462	AL033377	1.63
acid phosphatase, prostate	NM_001099	1.63
tripartite motif-containing 28	BE967532	1.63
signal sequence receptor, alpha (translocon-associated protein alpha)	NM_003144	1.63
hypothetical protein FLJ10043	NM_017979	1.63
Consensus includes gb:AI915947 /FEA=EST /DB_XREF=gi:5635802 /DB_XREF=est:wg96e01.x1 /CLONE=IMAGE:2379096 /UG=Hs.28212 ESTs	AI915947	1.63
crystallin, gamma B	NM_005210	1.63
disabled homolog 2, mitogen-responsive phosphoprotein (Drosophila)	N21202	1.63
Consensus includes gb:AK025029.1 /DEF=Homo sapiens cDNA: FLJ21376 fis, clone COL03231. /FEA=mRNA /DB_XREF=gi:10437464 /UG=Hs.306748 Homo sapiens cDNA: FLJ21376 fis, cl	AK025029	1.63
erythrocyte transmembrane protein	NM_016158	1.63
Consensus includes gb:AW970948 /FEA=EST /DB_XREF=gi:8160793 /DB_XREF=est:EST383031 /UG=Hs.269403 ESTs	AW970948	1.63
ets variant gene 1	X87175	1.63
hypothetical protein FLJ21162	NM_024873	1.63
chromosome 11open reading frame 9	AC004770	1.63

SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily c, member 1	AW152160	1.63
cysteine-rich motor neuron 1	BG546884	1.63
homeo box (H6 family) 1	NM_018942	1.63
bone morphogenetic protein 2	AA583044	1.63
hypothetical protein FLJ22686	AI825877	1.63
itchy homolog E3 ubiquitin protein ligase (mouse)	AL109923	1.63
epidermal growth factor receptor (erythroblastic leukemia viral (v-erb-b) oncogene homolog, avian)	AW157070	1.63
betacellulin	NM_001729	1.63
amyloid beta (A4) precursor protein-binding, family A, member 3 (X11-like 2)	NM_004886	1.63
Consensus includes gb:AI984051 /FEA=EST /DB_XREF=gi:5811270 /DB_XREF=est:wt52h03.x1 /CLONE=IMAGE:2511125 /UG=Hs.11861 thyroid hormone receptor-associated protein, 240	AI984051	1.63
hypothetical protein FLJ13480	NM_025120	1.63
nephronophthisis 1 (juvenile)	NM_000272	1.63
cadherin 15, M-cadherin (myotubule)	NM_004933	1.63
Consensus includes gb:AA830143 /FEA=EST /DB_XREF=gi:2903242 /DB_XREF=est:oc44e11.s1 /CLONE=IMAGE:1352588 /UG=Hs.15535 Homo sapiens clone 24582 mRNA sequence	AA830143	1.63
Kruppel-like factor 4 (gut)	BF514079	1.63
cadherin, EGF LAG seven-pass G-type receptor 1 (flamingo homolog, Drosophila)	NM_014246	1.63
hypothetical protein from BCRA2 region	NM_014887	1.63
adenosine monophosphate deaminase (isoform E)	AA919119	1.63
E2F-like protein	NM_016521	1.63
collagen, type III, alpha 1 (Ehlers-Danlos syndrome type IV, autosomal dominant)	AU144167	1.63
dystrobrevin, alpha	NM_001392	1.63
netrin 2-like (chicken)	NM_006181	1.63
6 antigen 1	NM_001468	1.63
vitelliform macular dystrophy (Best disease, bestrophin)	NM_004183	1.63
fibrinogen silencer binding protein	NM_006550	1.63
platelet-activating factor acetylhydrolase, isoform Ib, beta subunit (30kD)	BC000398	1.63
solute carrier family 4, sodium bicarbonate cotransporter, member 7	NM_003615	1.63
Oyster Ind. C18318:C18318 Homo sapiens cDNA, 5 end /clone=GEN-560E03 /clone_end=5 /gb=C18318 /gi=1579920 /ug=Hs.123469 /len=519	C18318	1.63
caveolin 1, caveolae protein, 22kD	NM_001753	1.63
protein tyrosine phosphatase, non-receptor type 12	S69182	1.63
relaxin 2 (H2)	NM_005059	1.63
FK506 binding protein 6 (36kD)	NM_003602	1.63
Consensus includes gb:AI742810 /FEA=EST /DB_XREF=gi:5111098 /DB_XREF=est:wg46e05.x1 /CLONE=IMAGE:2368160 /UG=Hs.131057 ESTs, Moderately similar to CRGD_HUMAN GAMM	AI742810	1.63
lipin 1	AA813260	1.63
Consensus includes gb:AL049798 /DEF=Human DNA sequence from clone 797M17 on chromosome 1q22-24.3. Contains the DPT gene for Dermatopontin, ESTs, an STS and GSSs /FEA=mRN	AL049798	1.63
hypothetical protein FLJ21628	NM_030613	1.63
choroideremia-like (Rab escort protein 2)	NM_001821	1.63
hypothetical protein DKFZp434H074	AL096732	1.63
Consensus includes gb:U79295.1 /DEF=Human clone 23961 mRNA sequence. /FEA=mRNA /DB_XREF=gi:1710277 /UG=Hs.90866 Human clone 23961 mRNA sequence	U79295	1.63
ets variant gene 6 (TEL oncogene)	AF041811	1.63
taste receptor, type 2, member 1	NM_019599	1.63
sushi-repeat-containing protein, X chromosome	NM_006307	1.63
STAT induced STAT inhibitor 3	BG035761	1.63
Mitochondrial Acyl-CoA Thioesterase	AF241787	1.63
Consensus includes gb:AK025564.1 /DEF=Homo sapiens cDNA: FLJ21911 fis, clone HEP03855. /FEA=mRNA /DB_XREF=gi:10438116 /UG=Hs.306816 Homo sapiens cDNA: FLJ21911 fis, cl	AK025564	1.63
hairy/enhancer-of-split related with YRPW motif 1	R61374	1.63
Consensus includes gb:AV718487 /FEA=EST /DB_XREF=gi:10815639 /DB_XREF=est:AV718487 /CLONE=GLCFVH09 /UG=Hs.282656 ESTs	AV718487	1.63
Consensus includes gb:U79304.1 /DEF=Human clone 23909 mRNA, partial cds. /FEA=mRNA /PROD=unknown /DB_XREF=gi:1710291 /UG=Hs.281043 Human clone 23909 mRNA, partial c	U79304	1.63
distal-less homeo box 6	NM_005222	1.63
transmembrane protein 5	NM_014254	1.63
KIAA0594 protein	AW183677	1.62
complement component (3b/4b) receptor 1, including Knops blood group system	X14362	1.62
hypothetical protein FLJ11191	NM_018355	1.62
carnitine palmitoyltransferase I, liver	BC000185	1.62
Consensus includes gb:AL030995 /DEF=Human DNA sequence from clone 1170D6 on chromosome Xq22.3-23. Contains a pseudogene similar to U-SNRNP_associated Cyclophilin (USA-CYP,	AL030995	1.62
myelin basic protein	NM_002385	1.62
glial cell derived neurotrophic factor	NM_000514	1.62
Consensus includes gb:AL023773 /DEF=Human DNA sequence from clone 406C18 on chromosome Xq27.1-27.3. Contains two MAGE (Melanoma Associated Antigen) family pseudogenes, GS	AL023773	1.62
erythrocyte membrane protein band 4.1-like 3	AI770004	1.62
Consensus includes gb:AF086790.1 /DEF=Homo sapiens aconitase precursor (ACON) mRNA, nuclear gene encoding mitochondrial protein, partial cds. /FEA=mRNA /GEN=ACON /PROD=aco	AF086790	1.62
lamin A/C	AA063189	1.62
hypothetical protein R29124_1	BF110434	1.62
gb:BC005220.1 /DEF=Homo sapiens, Similar to chaperonin containing TCP1, subunit 8 (theta), clone MGC:12240, mRNA, complete cds. /FEA=mRNA /PROD=Similar to chaperonin containin	BC005220	1.62

microrchidia homolog (mouse)	NM_014429	1.62
carbamoyl-phosphate synthetase 1, mitochondrial	AF154830	1.62
cysteine and histidine-rich domain (CHORD)-containing, zinc binding protein 1	NM_012124	1.62
lipidosin	AB014531	1.62
sialic acid binding Ig-like lectin 9	AF247180	1.62
Consensus includes gb:AL110190.1 /DEF=Homo sapiens mRNA; cDNA DKFZp564I2116 (from clone DKFZp564I2116). /FEA=mRNA /DB_XREF=gi:5817104 /UG=Hs.306338 Homo sapiens n	AL110190	1.62
glutamate receptor, metabotropic 5	NM_000842	1.62
serine (or cysteine) proteinase inhibitor, clade H (heat shock protein 47), member 1	NM_004353	1.62
golgi autoantigen, golgin subfamily a, 4	NM_002078	1.62
cytokine-inducible kinase	NM_004073	1.62
PRO1768 protein	NM_014099	1.62
testis expressed sequence 12	NM_031275	1.62
sulfotransferase family, cytosolic, 2B, member 1	NM_004605	1.62
Consensus includes gb:H49382 /FEA=EST /DB_XREF=gi:989223 /DB_XREF=est:yq19b06.s1 /CLONE=IMAGE:274115 /UG=Hs.124984 ESTs, Moderately similar to ALU7_HUMAN ALU SUBFA	H49382	1.62
Consensus includes gb:BF224076 /FEA=EST /DB_XREF=gi:11131303 /DB_XREF=est:7q83e08.x1 /CLONE=IMAGE:3704942 /UG=Hs.307438 Human DNA sequence from clone 495O10 on c	BF224076	1.62
SH3-domain binding protein 2	AB000462	1.62
alcohol dehydrogenase 1B (class I), beta polypeptide	M21692	1.62
hypothetical protein DKFZp434C0328	NM_017577	1.62
midline 1 (Opitz/BBB syndrome)	NM_000381	1.62
cytochrome P450, subfamily IIA (phenobarbital-inducible), polypeptide 7	NM_000764	1.62
zinc finger protein 144 (Mel-18)	BC004858	1.62
wingless-type MMTV integration site family, member 7B	BE736994	1.62
meprin A, beta	NM_005925	1.62
Consensus includes gb:U81961 /DEF=Homo sapiens amiloride-sensitive epithelial sodium channel alpha subunit alpha ENaC-1 and alpha ENaC-2 (SCNN1A) genes, alternatively spliced, prom	U81961	1.62
keratin 17	Z19574	1.62
mitogen-activated protein kinase-activated protein kinase 3	U43784	1.62
catenin (cadherin-associated protein), alpha 1 (102kD)	NM_001903	1.62
flotillin 1	AA507012	1.62
solute carrier family 22 (organic anion transporter), member 7	AF210455	1.62
N amino acid transporter 3	NM_018018	1.62
Consensus includes gb:AK022397.1 /DEF=Homo sapiens cDNA FLJ12335 fis, clone MAMMA1002219, highly similar to Rattus norvegicus rezo70 mRNA. /FEA=mRNA /DB_XREF=gi:1043378	AK022397	1.62
myosin XVA	NM_016239	1.62
distal-less homeobox 4	NM_001934	1.62
vanin 2	NM_004665	1.62
Consensus includes gb:AF070602.1 /DEF=Homo sapiens clone 24504 mRNA sequence. /FEA=mRNA /DB_XREF=gi:3387982 /UG=Hs.51649 Homo sapiens clone 24504 mRNA sequence	AF070602	1.62
SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily a, member 1	NM_003069	1.62
potassium inwardly-rectifying channel,subfamily J, member 14	NM_013348	1.62
keratin 18	AL031903	1.62
G protein-coupled receptor 23	NM_005296	1.62
CGI-62 protein	NM_016010	1.62
interleukin 9 receptor	L39064	1.62
KIAA1473 protein	AB040906	1.62
LIM protein (similar to rat protein kinase C-binding enigma)	BF671400	1.62
Consensus includes gb:AF070566.1 /DEF=Homo sapiens clones 24611 and 24705 mRNA sequence. /FEA=mRNA /DB_XREF=gi:3387934 /UG=Hs.302042 Homo sapiens clones 24611 and	AF070566	1.62
hypothetical protein MGC10946	NM_030572	1.62
nuclear factor I/B	BC001283	1.62
hypothetical protein FLJ20473	NM_017836	1.62
Consensus includes gb:AB051438.1 /DEF=Homo sapiens mRNA for KIAA1651 protein, partial cds. /FEA=mRNA /GEN=KIAA1651 /PROD=KIAA1651 protein /DB_XREF=gi:13359174 /UG=Hs.	AB051438	1.62
basic helix-loop-helix domain containing, class B, 2	NM_003670	1.62
histidine ammonia-lyase	NM_002108	1.62
nuclear receptor subfamily 1, group D, member 1	M24900	1.62
Consensus includes gb:AK024081.1 /DEF=Homo sapiens cDNA FLJ14019 fis, clone HEMBA1002503. /FEA=mRNA /DB_XREF=gi:10436370 /UG=Hs.287613 Homo sapiens cDNA FLJ14019 fi	AK024081	1.62
asporin (LRR class 1)	NM_017680	1.62
fibroblast growth factor 18	AB007422	1.62
collagen, type X, alpha 1 (Schmid metaphyseal chondrodysplasia)	AJ376003	1.62
protocadherin 1 (cadherin-like 1)	AA481656	1.62
surfactant, pulmonary-associated protein B	J02761	1.62
hypothetical protein PRO2796	NM_018633	1.62
hypothetical protein	NM_019099	1.62
fibroblast growth factor 5	AF171928	1.62
pregnancy-associated plasma protein-E	AF311940	1.62
cystein-rich hydrophobic domain 2	NM_012110	1.62
Consensus includes gb:BG230612 /FEA=EST /DB_XREF=gi:12725654 /DB_XREF=est:naf41b09.x1 /CLONE=IMAGE:4143329 /UG=Hs.13996 Homo sapiens cDNA: FLJ23260 fis, clone COL05	BG230612	1.62

melanocortin 4 receptor	NM_005912	1.62
RNA polymerase I transcription factor RRN3	AF001549	1.62
Kruppel-like factor 3 (basic)	NM_016531	1.62
solute carrier family 34 (sodium phosphate), member 1	NM_003052	1.62
transforming, acidic coiled-coil containing protein 2	NM_006997	1.62
immunoglobulin kappa constant	AF103574	1.62
selectin L (lymphocyte adhesion molecule 1)	NM_000655	1.62
chromosome 21 open reading frame 2	NM_004928	1.62
Consensus includes gb:R94644 /FEA=EST /DB_XREF=gi:970039 /DB_XREF=est:yq42a12.r1 /CLONE=IMAGE:198430 /UG=Hs.306542 Homo sapiens versican Vint isoform, mRNA, partial cd	R94644	1.62
hypothetical protein FLJ22792	NM_024921	1.61
Consensus includes gb:AW293343 /FEA=EST /DB_XREF=gi:6699905 /DB_XREF=est:UI-H-BI2-ahj-d-05-0-UI.s1 /CLONE=IMAGE:2726841 /UG=Hs.283978 Homo sapiens PRO2751 mRNA, c	AW293343	1.61
hypothetical protein DKFZp547H025	NM_020161	1.61
core 2 beta-1,6-N-acetylglucosaminyltransferase 3	NM_016591	1.61
potassium voltage-gated channel, subfamily H (eag-related), member 1	NM_002238	1.61
basement membrane-induced gene	NM_004848	1.61
sine oculis homeobox homolog 6 (Drosophila	NM_007374	1.61
hepatocyte growth factor (hepapoietin A; scatter factor)	U46010	1.61
disabled homolog 1 (Drosophila)	NM_021080	1.61
aryl hydrocarbon receptor nuclear translocator 2	NM_014862	1.61
HMP19 protein	NM_015980	1.61
dual oxidase 1	NM_017434	1.61
pregnancy-associated plasma protein A	NM_002581	1.61
aspartyl-tRNA synthetase	BC000629	1.61
retinoblastoma binding protein 9	NM_006606	1.61
jerky homolog-like (mouse)	NM_003772	1.61
ELAV (embryonic lethal, abnormal vision, Drosophila)-like 4 (Hu antigen D)	NM_021952	1.61
hypothetical protein	BC000039	1.61
phosphoinositide-3-kinase, catalytic, delta polypeptide	U57843	1.61
membrane-spanning 4-domains, subfamily A, member 2 (Fc fragment of IgE, high affinity I, receptor for; beta polypeptide)	D10583	1.61
serine (or cysteine) proteinase inhibitor, clade A (alpha-1 antiproteinase, antitrypsin), member 5	I02639	1.61
2.19 gene	BC002934	1.61
H4 histone family, member H	NM_003543	1.61
hypothetical gene FLJ00060	AK024467	1.61
potassium inwardly-rectifying channel, subfamily J, member 3	U39196	1.61
replication protein A complex 34 kd subunit homolog Rpa4	NM_013347	1.61
platelet activating receptor homolog	NM_013308	1.61
Consensus includes gb:X90579.1 /DEF=H.sapiens DNA for cyp related pseudogene. /FEA=mRNA /DB_XREF=gi:945005 /UG=Hs.166079 H.sapiens DNA for cyp related pseudogene	X90579	1.61
secreted frizzled-related protein 1	AF017987	1.61
Rho GDP dissociation inhibitor (GDI) gamma	NM_001176	1.61
interleukin 2 receptor, beta	NM_000878	1.61
phospholipase C, epsilon	NM_006226	1.61
adenylosuccinate lyase	AL022238	1.61
carbonic anhydrase I	NM_001738	1.61
tripartite motif-containing 29	AF230389	1.61
soluble adenylyl cyclase	NM_018417	1.61
Consensus includes gb:X90579.1 /DEF=H.sapiens DNA for cyp related pseudogene. /FEA=mRNA /DB_XREF=gi:945005 /UG=Hs.166079 H.sapiens DNA for cyp related pseudogene	X90579	1.61
Consensus includes gb:BG109855 /FEA=EST /DB_XREF=gi:12603361 /DB_XREF=est:602279521F1 /CLONE=IMAGE:4367308 /UG=Hs.49476 Homo sapiens clone TUA8 Cri-du-chat region	BG109855	1.61
regulated in glioma	NM_006394	1.61
hypothetical protein FLJ20043	NM_017637	1.61
thiopurine S-methyltransferase	BF196891	1.61
TI6 protein	AF112972	1.61
hypothetical protein FLJ10619	NM_018156	1.61
hypothetical protein FLJ20813	NM_017961	1.61
gb:U87229.1 /DEF=Homo sapiens death adaptor molecule RAIDD-2 mRNA, complete cds. /FEA=CDS /PROD=death adaptor molecule RAIDD-2 /DB_XREF=gi:5051867 /FL=gb:U87229.1	U87229	1.61
Consensus includes gb:AL359557.1 /DEF=Homo sapiens mRNA; cDNA DKFZp76201415 (from clone DKFZp76201415). /FEA=mRNA /DB_XREF=gi:8655612 /UG=Hs.306508 Homo sapiens	AL359557	1.61
fibulin 1	NM_006485	1.61
toll-like receptor 2	NM_003264	1.61
gamma-aminobutyric acid (GABA) A receptor, alpha 5	NM_000810	1.61
B-cell CLL/lymphoma 7A	AI950380	1.61
interleukin 3 (colony-stimulating factor, multiple)	NM_000588	1.61
dystrophin (muscular dystrophy, Duchenne and Becker types), includes DXS142, DXS164, DXS206, DXS230, DXS239, DXS268, DXS269, DXS270, DXS272	NM_004019	1.61
ER to nucleus signalling 2	AI732416	1.61
cell growth regulatory with EF-hand domain	NM_006569	1.61

crystallin, beta B1	NM_001887	1.61
synovial sarcoma, X breakpoint 3	BF184274	1.61
Consensus includes gb:X69383 /DEF=H.sapiens gene for T cell receptor gamma V region 5 /FEA=CDS_2 /DB_XREF=gi:37009 /UG=Hs.247801 H.sapiens gene for T cell receptor gamma V r	X69383	1.61
adrenergic, beta, receptor kinase 1	NM_001619	1.61
potassium voltage-gated channel, shaker-related subfamily, member 2	NM_004974	1.61
phosphatidylinositol-4-phosphate 5-kinase, type I, beta	NM_003558	1.61
actinin, alpha 2	W63731	1.61
S100 calcium binding protein A9 (calgranulin B)	NM_002965	1.61
lymphocyte antigen 94 homolog, activating NK-receptor; NK-p46, (mouse)	AJ006122	1.61
Consensus includes gb:AF052140.1 /DEF=Homo sapiens clone 23758 mRNA sequence. /FEA=mRNA /DB_XREF=gi:3360449 /UG=Hs.141055 Homo sapiens clone 23758 mRNA sequence	AF052140	1.61
hypothetical protein MGC5178	AI580112	1.61
glucagon	NM_002054	1.61
Mahlavu hepatocellular carcinoma	NM_006543	1.61
ectonucleoside triphosphate diphosphohydrolase 1	NM_001776	1.61
Consensus includes gb:AW613387 /FEA=EST /DB_XREF=gi 7318573 /DB_XREF=est:hh71e04.x1 /CLONE=IMAGE:2968254 /UG=Hs.278541 ESTs, Moderately similar to TYPH_HUMAN THYR	AW613387	1.61
nuclear receptor subfamily 1, group I, member 3	NM_005122	1.61
G-protein coupled receptor	NM_007369	1.61
pre-B-cell leukemia transcription factor 2	BE397715	1.61
musculin (activated B-cell factor-1)	AF060154	1.61
glucan (1,4-alpha-), branching enzyme 1 (glycogen branching enzyme, Andersen disease, glycogen storage disease type IV)	NM_000158	1.61
N-acylsphingosine amidohydrolase (acid ceramidase)	AI379338	1.61
HSPC039 protein	AF119875	1.61
PTEN induced putative kinase 1	BF432478	1.61
uncharacterized bone marrow protein BM029	AI679080	1.61
chromosome 12 open reading frame 6	NM_020367	1.61
pvt-1 (murine) oncogene homolog, MYC activator	M34428	1.61
FXYD domain-containing ion transport regulator 2	NM_001680	1.61
paraneoplastic antigen MA2	AF286487	1.61
DnaJ (Hsp40) homolog, subfamily C, member 8	AF116696	1.61
Consensus includes gb:AF052090.1 /DEF=Homo sapiens clone 23950 mRNA sequence. /FEA=mRNA /DB_XREF=gi:3360396 /UG=Hs.106620 Homo sapiens clone 23950 mRNA sequence	AF052090	1.61
epidermal growth factor receptor substrate EPS15R	AV710549	1.61
hypothetical protein	AK026747	1.61
MCF.2 cell line derived transforming sequence	NM_005369	1.61
cadherin 11, type 2, OB-cadherin (osteoblast)	D21254	1.61
HDCMC28P protein	NM_016649	1.61
Consensus includes gb:AL080232.1 /DEF=Homo sapiens mRNA; cDNA DKFZp586A061 (from clone DKFZp586A061). /FEA=mRNA /DB_XREF=gi:5262725 /UG=Hs.220696 Homo sapiens mR	AL080232	1.61
paired immunoglobulin-like receptor alpha	NM_013439	1.61
transferrin	AI073407	1.61
LAP (leucine-rich repeats and PDZ) and no PDZ protein	NM_025168	1.61
insulin-like 3 (Leydig cell)	NM_005543	1.61
hypothetical protein FLJ21919	AF130066	1.61
myomesin (M-protein) 2 (165kD)	NM_003970	1.60
Bcl-2 binding component 3	AF332558	1.60
receptor tyrosine kinase-like orphan receptor 2	NM_004560	1.60
insulinoma-associated 1	NM_002196	1.60
Consensus includes gb:AL390183.1 /DEF=Homo sapiens mRNA; cDNA DKFZp547D096 (from clone DKFZp547D096). /FEA=mRNA /DB_XREF=gi:9368895 /UG=Hs.88121 Homo sapiens mRI	AL390183	1.60
EGF-like repeats and discoidin I-like domains 3	NM_005711	1.60
KIAA0296 gene product	NM_014699	1.60
Consensus includes gb:U66046.1 /DEF=Homo sapiens clone 161455 breast expressed mRNA from chromosome X. /FEA=mRNA /DB_XREF=gi:1519271 /UG=Hs.92683 Homo sapiens clone	U66046	1.60
phospholipid scramblase 4	NM_020353	1.60
CGI-119 protein	NM_016056	1.60
SH3-domain protein 5 (ponsin)	NM_015385	1.60
hypothetical protein FLJ20313	NM_017762	1.60
Consensus includes gb:AL136548.1 /DEF=Homo sapiens mRNA; cDNA DKFZp761G18121 (from clone DKFZp761G18121); complete cds. /FEA=mRNA /GEN=DKFZp761G18121 /PROD=hypc	AL136548	1.60
HSPC135 protein	NM_014170	1.60
hypothetical protein LOC57333	AI797684	1.60
hypothetical protein FLJ14126	NM_024849	1.60
KIAA1025 protein	BG426689	1.60
activating transcription factor 5	AF101388	1.60
N-deacetylase/N-sulfotransferase (heparan glucosaminyl) 1	AL526632	1.60
chemokine binding protein 2	NM_001296	1.60
immunoglobulin kappa constant	AF103530	1.60
p21 (CDKN1A)-activated kinase 3	AW085556	1.60

FE65-LIKE 2	NM_006051	1.60
phosphofipase C, delta 1	NM_006225	1.60
DKFZP566K0524 protein	AL050040	1.60
collagen, type II, alpha 1 (primary osteoarthritis, spondyloepiphyseal dysplasia, congenital)	X16468	1.60
Consensus includes gb:AW301775 /FEA=EST /DB_XREF=gi:6711452 /DB_XREF=est:zr83e06.z1 /CLONE=IMAGE:2766754 /UG=Hs.254727 ESTs	AW301775	1.60
a disintegrin and metalloproteinase domain 22	AF155381	1.60
empty spiracles homolog 1 (Drosophila)	BF515060	1.60
v-Ki-ras2 Kirsten rat sarcoma 2 viral oncogene homolog	BF673699	1.60
hypothetical protein FLJ10210	NM_018027	1.60
hypothetical protein FLJ10244	NM_018037	1.60
pregnancy-associated plasma protein A	L13197	1.60
ORF	BC001372	1.60
Usher syndrome 2A (autosomal recessive, mild)	NM_007123	1.60
Wiskott-Aldrich syndrome (eczema-thrombocytopenia)	NM_000377	1.60
glutathione peroxidase 3 (plasma)	NM_002084	1.60
development and differentiation enhancing factor 1	NM_018482	1.60
hypothetical protein FLJ23047	NM_024548	1.60
Consensus includes gb:AL049244.1 /DEF=Homo sapiens mRNA; cDNA DKFZp564C163 (from clone DKFZp564C163). /FEA=mRNA /DB_XREF=gi:4499979 /UG=Hs.302046 Homo sapiens mR	AL049244	1.60
polymerase (RNA) II (DNA directed) polypeptide J (13.3kD)	BE676209	1.60
adaptor-related protein complex 1, mu 2 subunit	NM_005498	1.60
carnitine palmitoyltransferase I, liver	BF001714	1.60
1-acylglycerol-3-phosphate O-acyltransferase 3	NM_020132	1.60
6-phosphofructo-2-kinase/fructose-2,6-biphosphatase 4	NM_004567	1.60
Src-like-adaptor	NM_006748	1.60
Dombrock blood group	NM_021071	1.60
Consensus includes gb:BE877397 /FEA=EST /DB_XREF=gi:10326173 /DB_XREF=est:601485434F1 /CLONE=IMAGE:3887936 /UG=Hs.240140 Human DNA sequence from clone 153G14 on	BE877397	1.60
fucosyltransferase 5 (alpha (1,3) fucosyltransferase)	U27329	1.60
gb:BC001217.1 /DEF=Homo sapiens, Similar to KIAA0029 protein, clone MGC:767, mRNA, complete cds. /FEA=mRNA /PROD=Similar to KIAA0029 protein /DB_XREF=gi:12654748 /UG=Hs.	BC001217	1.60
kelch-like 2, Mayven (Drosophila)	NM_007246	1.60
Consensus includes gb:AK025143.1 /DEF=Homo sapiens cDNA: FLJ21490 fis, clone COL05464. /FEA=mRNA /DB_XREF=gi:10437602 /UG=Hs.288700 Homo sapiens cDNA: FLJ21490 fis, cl	AK025143	1.60
wingless-type MMTV integration site family, member 8B	NM_003393	1.60
cadherin 10, type 2 (T2-cadherin)	NM_006727	1.60
Consensus includes gb:BG391282 /FEA=EST /DB_XREF=gi:13284730 /DB_XREF=est:602417327F1 /CLONE=IMAGE:4536582 /UG=Hs.306000 hypothetical protein FLJ10624	BG391282	1.60
regulator of G-protein signalling 6	AF156932	1.60
leukocyte immunoglobulin-like receptor, subfamily A (with TM domain), member 2	U82277	1.60
death-associated protein kinase 1	BC003614	1.60
Consensus includes gb:D29810.1 /DEF=Human mRNA for unknown product, partial cds. /FEA=mRNA /PROD=unknown /DB_XREF=gi:704440 /UG=Hs.153445 Human mRNA for unknown pr	D29810	1.60
ATPase, Class V, type 10C	AB011138	1.60
hypothetical protein FLJ10718	NM_018192	1.60
NCK protein	NM_016170	1.60
KIAA1024 protein	AF035294	1.60
hypothetical protein MGC5338	NM_024062	1.60
leptin receptor	NM_002303	1.60
MAX dimerization protein	NM_002357	1.60
plastin 1 (I isoform)	NM_002670	1.60
matrix metalloproteinase 14 (membrane-inserted)	AU149305	1.60
Cluster Incl. N32185:yy25d10.s1 Homo sapiens cDNA, 3' end /clone=IMAGE-272275 /clone_end=3' /gb=N32185 /gi=1152584 /ug=Hs.120769 /len=600	N32185	1.60
ring finger protein (C3H2C3 type) 6	AF293342	1.60
thyroid stimulating hormone receptor	BE045816	1.60
mesenchymal stem cell protein DSC43	AF242768	1.60
cadherin 6, type 2, K-cadherin (fetal kidney)	BC000019	1.60
hypothetical protein FLJ22843	NM_025184	1.60
hypothetical protein FLJ11328	NM_018391	1.60
hypothetical protein FLJ13385	NM_024853	1.60
dystrobrevin, beta	Y15718	1.60
pp32 related 1	NM_012403	1.60
ribosome binding protein 1 homolog 180kD (dog)	AA706065	1.60
lymphocyte antigen 117	AF000426	1.60
hypothetical protein FLJ10770	NM_018211	1.60
GDNF family receptor alpha 3	NM_001496	1.60
kruppel-related zinc finger protein hcKrox	NM_015872	1.60
Consensus includes gb:AK022421.1 /DEF=Homo sapiens cDNA FLJ12359 fis, clone MAMMA1002355. /FEA=mRNA /DB_XREF=gi:10433814 /UG=Hs.296718 Homo sapiens cDNA FLJ12359	AK022421	1.60
Kruppel-type zinc finger (C2H2)	NM_005815	1.60

hypothetical protein FLJ10786	NM_018219	1.60
solute carrier family 16 (monocarboxylic acid transporters), member 7	AF049608	1.60
ATP-binding cassette, sub-family B (MDR/TAP), member 9	BE504895	1.60
filamin C, gamma (actin binding protein 280)	NM_001458	1.60
Consensus includes gb:AU144900 /FEA=EST /DB_XREF=gi:11006421 /DB_XREF=est:AU144900 /CLONE=HEMBA1003327 /UG=Hs.274233 Homo sapiens cDNA FLJ10146 fis, clone HEMBA	AU144900	1.60
slug homolog, zinc finger protein (chicken)	AI572079	1.60
hypothetical protein MGC2742	NM_023938	1.60
VLCS-H1 protein	NM_014031	1.60
engulfment and cell motility 1 (ced-12 homolog, C. elegans)	NM_014800	1.60
Consensus includes gb:AK023959.1 /DEF=Homo sapiens cDNA FLJ13897 fis, clone THYRO1001706. /FEA=mRNA /DB_XREF=gi:10436079 /UG=Hs.301488 Homo sapiens cDNA FLJ13897 fi	AK023959	1.60
FOS-like antigen 2	N36408	1.60
hypothetical protein FLJ22344	NM_024717	1.59
cytochrome b-561	U06715	1.59
SEC10-like 1 (S. cerevisiae)	NM_006544	1.59
RAP2B, member of RAS oncogene family	AW005535	1.59
hypothetical protein FLJ20502	NM_017845	1.59
hypothetical protein PRO2176	NM_018515	1.59
toll-like receptor 6	NM_006068	1.59
mucin 1, transmembrane	AI610869	1.59
hypothetical protein FLJ23554	NM_024806	1.59
KIAA0485 protein	AW129056	1.59
suc1-associated neurotrophic factor target (FGFR signalling adaptor)	NM_006654	1.59
Consensus includes gb:AI480014 /FEA=EST /DB_XREF=gi:4373182 /DB_XREF=est:tm71b09.x1 /CLONE=IMAGE:2163545 /UG=Hs.13415 Homo sapiens clone 24571 mRNA sequence	AI480014	1.59
potassium voltage-gated channel, Shab-related subfamily, member 2	AF338730	1.59
protein tyrosine phosphatase type IVA, member 3	BC003105	1.59
Consensus includes gb:AL551046 /FEA=EST /DB_XREF=gi:12888616 /DB_XREF=est:AL551046 /CLONE=CS0DI066YN13 (5 prime) /UG=Hs.332742 Human DNA from chromosome 19-spec	AL551046	1.59
B-cell CLL/lymphoma 2	NM_000633	1.59
protease, serine, 23	NM_007173	1.59
interleukin 1 receptor antagonist	U65590	1.59
hemochromatosis	NM_000410	1.59
Consensus includes gb:AL080104.1 /DEF=Homo sapiens mRNA; cDNA DKFZp566D153 (from clone DKFZp566D153). /FEA=mRNA /DB_XREF=gi:5262529 /UG=Hs 306322 Homo sapiens mI	AL080104	1.59
Cdc42 effector protein 4; binder of Rho GTPases 4	AF099664	1.59
protocadherin beta 8	NM_019120	1.59
eukaryotic translation initiation factor 3, subunit 10 (theta, 150/170kD)	NM_003750	1.59
microfibrillar-associated protein 3	R38475	1.59
gb:L14453.1 /DEF=Human Ig rearranged mu-chain gene V-N-D-N-J-region, complete cds. /FEA=mRNA /DB_XREF=gi:398494 /FL=gb:L14453.1	L14453	1.59
RAB-8b protein	NM_016530	1.59
peptidyl arginine deiminase, type V	AF229067	1.59
hypothetical protein FLJ21369	NM_024802	1.59
NK-receptor	AF022048	1.59
Consensus includes gb:AK024851.1 /DEF=Homo sapiens cDNA: FLJ21198 fis, clone COL00220. /FEA=mRNA /DB_XREF=gi:10437259 /UG=Hs.306707 Homo sapiens cDNA: FLJ21198 fis, cl	AK024851	1.59
arsenate resistance protein ARS2	H28020	1.59
peptidylprolyl isomerase E (cyclophilin E)	AF104013	1.59
transmembrane 4 superfamily member 1	AI189753	1.59
leukocyte tyrosine kinase	NM_002344	1.59
Consensus includes gb:AF007137.1 /DEF=Homo sapiens clone 23618 mRNA sequence. /FEA=mRNA /DB_XREF=gi:2852612 /UG=Hs.302185 Homo sapiens clone 23618 mRNA sequence	AF007137	1.59
Consensus includes gb:AL117652.1 /DEF=Homo sapiens mRNA; cDNA DKFZp586B1324 (from clone DKFZp586B1324). /FEA=mRNA /DB_XREF=gi:5912244 /UG=Hs.278334 Homo sapiens	AL117652	1.59
colony stimulating factor 3 (granulocyte)	NM_000759	1.59
neuropilin 1	BE620457	1.59
testis expressed sequence 11	NM_031276	1.59
Consensus includes gb:AW972765 /FEA=EST /DB_XREF=gi:8162611 /DB_XREF=est:EST384859 /UG=Hs.293299 ESTs	AW972765	1.59
Consensus includes gb:AU152579 /FEA=EST /DB_XREF=gi:11014100 /DB_XREF=est:AU152579 /CLONE=NT2RP3001232 /UG=Hs.288931 Homo sapiens cDNA FLJ13034 fis, clone NT2RP3	AU152579	1.59
Bacillus subtilis /REF=L38424 /DEF=B subtilis dapB, jojF, jojG genes corresponding to nucleotides 2055-2578 of L38424 /LEN=1931 (-5, -M, -3 represent transcript regions 5 prime, Midd	L38424	1.59
hypothetical protein FLJ12345	NM_024987	1.59
von Hippel-Lindau syndrome	NM_000551	1.59
zinc finger protein 185 (LIM domain)	NM_007150	1.59
CD58 antigen, (lymphocyte function-associated antigen 3)	BC005930	1.59
Consensus includes gb:AL571723 /FEA=EST /DB_XREF=gi:12929303 /DB_XREF=est:AL571723 /CLONE=CS0DI031YD10 (3 prime) /UG=Hs.7247 ESTs, Weakly similar to TIE1_HUMAN TYR(	AL571723	1.59
MG36 protein	NM_025256	1.59
lectomedin-2	NM_014921	1.59
hypothetical protein	NM_016449	1.59
Mdm2, transformed 3T3 cell double minute 2, p53 binding protein (mouse)	AF201370	1.59
hypothetical protein FLJ10300	NM_018051	1.59

Consensus includes gb:M31949 /DEF=Human Ig rearranged mu-chain V-region gene, subgroup VH-III, exon 1 and 2 /FEA=CDS /DB_XREF=gi:185254 /UG=Hs.247923 Human Ig rearranged	M31949	1.59
G1 to S phase transition 2	NM_018094	1.59
tumor-associated calcium signal transducer 2	AI627697	1.59
KIAA1077 protein	BE500977	1.59
Consensus includes gb:R71245 /FEA=EST /DB_XREF=gi:844762 /DB_XREF=est:yi54e05.s1 /CLONE=IMAGE:143072 /UG=Hs.174303 ESTs	R71245	1.59
tubby homolog (mouse)	U82467	1.59
hypothetical protein FLJ11800	NM_024974	1.59
synapsin I	H19843	1.59
B-cell CLL/lymphoma 7A	BC002629	1.59
cadherin 20, type 2	AF217289	1.59
Consensus includes gb:AK022038.1 /DEF=Homo sapiens cDNA FLJ11976 fis, clone HEMBB1001253. /FEA=mRNA /DB_XREF=gi:10433356 /UG=Hs.296679 Homo sapiens cDNA FLJ11976 f	AK022038	1.59
KIAA0944 protein	AB023161	1.59
FKSG17	AF315951	1.59
growth factor receptor-bound protein 10	U66065	1.59
Consensus includes gb:AK024185.1 /DEF=Homo sapiens cDNA FLJ14123 fis, clone MAMMA1002155. /FEA=mRNA /DB_XREF=gi:10436502 /UG=Hs.269314 Homo sapiens cDNA FLJ14123 f	AK024185	1.59
Consensus includes gb:AL050145.1 /DEF=Homo sapiens mRNA; cDNA DKFZp586C2020 (from clone DKFZp586C2020). /FEA=mRNA /DB_XREF=gi:4884356 /UG=Hs.225986 Homo sapiens	AL050145	1.59
short stature homeobox 2	AI816713	1.59
KIAA0754 protein	AW663712	1.59
transmembrane protein with EGF-like and two follistatin-like domains 1	BF439316	1.59
interferon, alpha-inducible protein 27	NM_005532	1.59
KIAA0942 protein	AW117368	1.59
homeo box D11	NM_021192	1.59
Rho guanine exchange factor (GEF) 12	NM_015313	1.59
Consensus includes gb:AU150943 /FEA=EST /DB_XREF=gi:11012464 /DB_XREF=est:AU150943 /CLONE=NT2RP2003984 /UG=Hs.66762 Homo sapiens mRNA; cDNA DKFZp564A026 (from	AU150943	1.59
glial fibrillary acidic protein	NM_002055	1.59
ATPase, Ca++ transporting, plasma membrane 3	AW615612	1.59
hypothetical protein FLJ22282	NM_024792	1.59
homeo box B6	AA527340	1.59
NADH:ubiquinone oxidoreductase MLRQ subunit homolog	NM_020142	1.59
sialyltransferase 4A (beta-galactosidase alpha-2,3-sialytransferase)	NM_003033	1.59
chymotrypsinogen B1	NM_001906	1.59
acyl-Coenzyme A dehydrogenase, C-2 to C-3 short chain	NM_000017	1.59
ficolin (collagen/fibrinogen domain-containing lectin) 2 (hucolin)	NM_015839	1.59
lactalbumin, alpha-	NM_002289	1.59
KIAA0308 protein	BF668950	1.59
gb:M26463.1 /DEF=Homo sapiens immunoglobulin mu chain antibody M030 (IgM) mRNA, complete cds. /FEA=CDS /GEN=IgM /PROD=immunoglobulin mu chain antibody M030 /DB_XREF=	M26463	1.59
hect domain and RLD 3	NM_014606	1.59
neurexin 2	AA608820	1.59
phosphorylase kinase, gamma 1 (muscle)	NM_006213	1.59
cartilage intermediate layer protein, nucleotide pyrophosphohydrolase	NM_003613	1.59
BAI1-associated protein 2	BC002495	1.59
Consensus includes gb:AL139318 /DEF=Human DNA sequence from clone RP11-124B17 on chromosome 13. Contains the DCT gene for dopachrome tautomerase (dopachrome delta-isomer	AL139318	1.59
Consensus includes gb:AL049767 /DEF=Human DNA sequence from clone 172H20 on chromosome 20q12-13.12 Contains genes for semenogelin I and II and PI3 (protease inhibitor 3, skin-	AL049767	1.59
hypothetical protein FLJ21065	NM_022481	1.59
hypothetical protein from clone 24796	AF070596	1.59
2'-5'-oligoadenylate synthetase-like	NM_003733	1.59
neurotrophic tyrosine kinase, receptor, type 3	S76476	1.59
piccolo (presynaptic cytomatrix protein)	AB011131	1.59
calpain 6	AI000388	1.59
myeloperoxidase	J02694	1.59
myeloid/lymphoid or mixed-lineage leukemia (trithorax homolog, Drosophila); translocated to, 4	AB016902	1.59
dual adaptor of phosphotyrosine and 3-phosphoinositides	NM_014395	1.59
Consensus includes gb:AF038181.1 /DEF=Homo sapiens clone 23607 mRNA sequence. /FEA=mRNA /DB_XREF=gi:2795901 /UG=Hs.167825 Homo sapiens clone 23607 mRNA sequence	AF038181	1.59
Consensus includes gb:AK001442.1 /DEF=Homo sapiens cDNA FLJ10580 fis, clone NT2RP2003533. /FEA=mRNA /DB_XREF=gi:7022701 /UG=Hs.140402 Homo sapiens cDNA FLJ10580 fis	AK001442	1.59
retinoid X receptor, beta	BF337038	1.59
glycophorin A (includes MN blood group)	NM_002099	1.59
KIAA1922 protein	BE300056	1.59
retinoblastoma binding protein 4	AL390237	1.59
NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 2 (8kD, B8)	AA993683	1.58
growth arrest and DNA-damage-inducible, beta	AV658684	1.58
antigen identified by monoclonal antibody MRC OX-2	AF063591	1.58
v-myc myelocytomatosis viral related oncogene, neuroblastoma derived (avian)	BC002712	1.58
galanin	AL556409	1.58

protein phosphatase 1, regulatory (inhibitor) subunit 12B	NM_002481	1.58
phosphodiesterase 1A, calmodulin-dependent	NM_005019	1.58
acid-inducible phosphoprotein	NM_006107	1.58
translocase of inner mitochondrial membrane 17 homolog A (yeast)	AK023063	1.58
karyopherin beta 2b, transportin	AK022217	1.58
calpastatin	D16217	1.58
gamma-aminobutyric acid (GABA) A receptor, pi	NM_014211	1.58
hypothetical protein FLJ11996	NM_024976	1.58
chromosome 1 open reading frame 10	NM_016190	1.58
thymosin, beta, identified in neuroblastoma cells	BF677486	1.58
zeta-chain (TCR) associated protein kinase (70 kD)	AI817942	1.58
frizzled-related protein	U91903	1.58
early lymphoid activation protein	L22650	1.58
hypothetical protein DKFZp761F2014	NM_020215	1.58
Consensus includes gb:AW675473 /FEA=EST /DB_XREF=gi:7540699 /DB_XREF=est:ba49g05.x1 /CLONE=IMAGE:2899928 /UG=Hs.101916 Homo sapiens mRNA; cDNA DKFZp564K133 (fro	AW675473	1.58
Consensus includes gb:M14087.1 /DEF=Human HL14 gene encoding beta-galactoside-binding lectin, 3 end, clone 2. /FEA=mRNA /DB_XREF=gi:187113 /UG=Hs.287389 Human HL14 gen	M14087	1.58
phosphodiesterase 4D, cAMP-specific (phosphodiesterase E3 dunce homolog, Drosophila)	AF012073	1.58
lethal (3) malignant brain tumor l(3)mbt protein (Drosophila) homolog	NM_015478	1.58
gb:M86841.1 /DEF=Human serotonin receptor type 2 (5HT2) mRNA, complete cds. /FEA=mRNA /GEN=5HT2 /PROD=serotonin receptor /DB_XREF=gi:177775 /FL=gb:M86841.1	M86841	1.58
brain-derived neurotrophic factor	NM_001709	1.58
thyroid hormone receptor interactor 6	AF000974	1.58
RAD51-like 1 (S. cerevisiae)	U84138	1.58
KIAA0665 gene product	NM_014700	1.58
glycine amidinotransferase (L-arginine:glycine amidinotransferase)	X86401	1.58
Ewing sarcoma breakpoint region 1	BC000527	1.58
Consensus includes gb:AI078169 /FEA=EST /DB_XREF=gi:3412577 /DB_XREF=est:oz30d11.x1 /CLONE=IMAGE:1676853 /UG=Hs.323511 Homo sapiens cDNA: FLJ23176 fis, clone LNG10	AI078169	1.58
protein phosphatase 3 (formerly 2B), catalytic subunit, alpha isoform (calcineurin A alpha)	NM_000944	1.58
catenin (cadherin-associated protein), beta 1 (88kD)	NM_001904	1.58
scrapie responsive protein 1	NM_007281	1.58
dual specificity phosphatase 4	NM_001394	1.58
flavin containing monooxygenase 2	AL021026	1.58
hypothetical protein LOC57821	NM_021179	1.58
ectodermal dysplasia 1, anhidrotic	AF061192	1.58
phospholipase A2, group IIA (platelets, synovial fluid)	NM_000300	1.58
Consensus includes gb:AF090886.1 /DEF=Homo sapiens clone HQ0072. /FEA=mRNA /DB_XREF=gi:6690153 /UG=Hs.284192 Homo sapiens clone HQ0072	AF090886	1.58
putative ankyrin-repeat containing protein	NM_015623	1.58
paraoxonase 1	NM_000446	1.58
adenosine A1 receptor	X68485	1.58
keratin 6B	AI831452	1.58
3-phosphoinositide dependent protein kinase-1	NM_002613	1.58
Consensus includes gb:BF002254 /FEA=EST /DB_XREF=gi:10702529 /DB_XREF=est:7h02a03.x1 /CLONE=IMAGE:3314764 /UG=Hs.143600 type II Golgi membrane protein /FL=gb:U5585	BF002254	1.58
transient receptor potential channel 3	U47050	1.58
carcinoembryonic antigen-related cell adhesion molecule 5	NM_004363	1.58
hypothetical protein FLJ23506	NM_024833	1.58
exportin, tRNA (nuclear export receptor for tRNAs)	AI984005	1.58
HGC6.2 protein	NM_014356	1.58
hypothetical protein FLJ14298	NM_024764	1.58
Consensus includes gb:AV651668 /FEA=EST /DB_XREF=gi:9872682 /DB_XREF=est:AV651668 /CLONE=GLCCSC04 /UG=Hs.282480 ESTs	AV651668	1.58
Consensus includes gb:AV720514 /FEA=EST /DB_XREF=gi:10817666 /DB_XREF=est:AV720514 /CLONE=GLCGSB09 /UG=Hs.282721 ESTs, Weakly similar to ALU7_HUMAN ALU SUBFAMILY	AV720514	1.58
hypothetical protein FLJ22955	NM_024819	1.58
DKFZP564A122 protein	AA521269	1.58
Consensus includes gb:AK023918.1 /DEF=Homo sapiens cDNA FLJ13856 fis, clone THYRO1000988. /FEA=mRNA /DB_XREF=gi:10436003 /UG=Hs.288489 Homo sapiens cDNA FLJ13856 fi	AK023918	1.58
nucleolar protein 3 (apoptosis repressor with CARD domain)	AF043244	1.58
nescient helix loop helix 1	H96739	1.58
PR domain containing 11	NM_020229	1.58
fibroblast growth factor 2 (basic)	NM_002006	1.58
matrix metalloproteinase 24 (membrane-inserted)	NM_006690	1.58
hypothetical protein FLJ10922	NM_018273	1.58
CGI-86 protein	AW243128	1.58
AXL receptor tyrosine kinase	NM_021913	1.58
nuclear fragile X mental retardation protein interacting protein 1	NM_012345	1.58
protamine 2	BC005303	1.58
lymphoid blast crisis oncogene	NM_006738	1.58

cytochrome P450, subfamily IIB (phenobarbital-inducible), polypeptide 6	X06399	1.58
CASP8 and FADD-like apoptosis regulator	AF015452	1.58
Consensus includes gb:AW134608 /FEA=EST /DB_XREF=gi:6138154 /DB_XREF=est:UI-H-BI1-abo-e-04-0-UI.s1 /CLONE=IMAGE:2712654 /UG=Hs.168941 ESTs	AW134608	1.58
Consensus includes gb:AK024938.1 /DEF=Homo sapiens cDNA: FLJ21285 fis, clone COL01912. /FEA=mRNA /DB_XREF=gi:10437359 /UG=Hs.287655 Homo sapiens cDNA: FLJ21285 fis, cl	AK024938	1.58
hypothetical protein MGC4293	BC002817	1.58
solute carrier family 15 (oligopeptide transporter), member 1	AB001328	1.58
CDP-diacylglycerol synthase (phosphatidate cytidylyltransferase) 1	NM_001263	1.58
H2A histone family, member X	BE901081	1.58
gb:NM_006273.2 /DEF=Homo sapiens small inducible cytokine A7 (monocyte chemotactic protein 3) (SCYA7), mRNA. /FEA=mRNA /GEN=SCYA7 /PROD=monocyte chemotactic protein 3 pr	NM_006273	1.58
phosphatidylinositol-4-phosphate 5-kinase, type I, alpha	U78576	1.58
zinc finger protein 131 (clone pHZ-10)	BE972394	1.58
adrenergic, beta-1-, receptor	NM_000684	1.58
synuclein, alpha interacting protein (synphilin)	NM_005460	1.58
hyaluronoglucosaminidase 1	AF173154	1.58
hypothetical protein FLJ22419	NM_024697	1.58
CREB binding protein (Rubinstein-Taybi syndrome)	U85962	1.58
Consensus includes gb:AK022442.1 /DEF=Homo sapiens cDNA FLJ12380 fis, clone MAMMA1002556. /FEA=mRNA /DB_XREF=gi:10433838 /UG=Hs.300934 Homo sapiens cDNA FLJ12380 f	AK022442	1.58
gb:BC002778.1 /DEF=Homo sapiens, Similar to myosin light chain 2, precursor lymphocyte-specific, clone MGC:3479, mRNA, complete cds. /FEA=mRNA /PROD=Similar to myosin light chai	BC002778	1.58
annexin A10	AF196478	1.58
gap junction protein, alpha 8, 50kD (connexin 50)	NM_005267	1.58
Consensus includes gb:AC004010 /DEF=Human BAC clone GS1-99H8 /FEA=CDS /DB_XREF=gi:2781385 /UG=Hs.121520 Human BAC clone GS1-99H8	AC004010	1.58
glutamate receptor, ionotropic, kainate 5	AL137705	1.58
potassium voltage-gated channel, Shab-related subfamily, member 2	NM_004770	1.58
hypothetical protein	NM_018836	1.57
integral membrane protein 2B	NM_021999	1.57
epithelial V-like antigen 1	AF275945	1.57
erythrocyte membrane protein band 4.1-like 3	BC006141	1.57
old astrocyte specifically induced substance	BG328407	1.57
Consensus includes gb:AK024393.1 /DEF=Homo sapiens cDNA FLJ14331 fis, clone PLACE4000320. /FEA=mRNA /DB_XREF=gi:10436771 /UG=Hs.288569 Homo sapiens cDNA FLJ14331 fi	AK024393	1.57
interleukin 1 receptor accessory protein-like 2	NM_017416	1.57
Consensus includes gb:AK025174.1 /DEF=Homo sapiens cDNA: FLJ21521 fis, clone COL05880. /FEA=mRNA /DB_XREF=gi:10437636 /UG=Hs.306777 Homo sapiens cDNA: FLJ21521 fis, cl	AK025174	1.57
hypothetical protein FLJ22596	NM_025086	1.57
calcium channel, voltage-dependent, alpha 1E subunit	NM_000721	1.57
gamma-aminobutyric acid (GABA) A receptor, alpha 2	NM_000807	1.57
Consensus includes gb:AL117520.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434P2435 (from clone DKFZp434P2435). /FEA=mRNA /DB_XREF=gi:5912036 /UG=Hs.306348 Homo sapiens	AL117520	1.57
LIM homeobox protein 2	NM_004789	1.57
POM (POM121 rat homolog) and ZP3 fusion	BC000487	1.57
HIV-1 Tat interactive protein 2, 30 kDa	AF092095	1.57
hypothetical protein FLJ23560	NM_024685	1.57
cholesteryl ester transfer protein, plasma	NM_000078	1.57
chromosome 15 open reading frame 5	NM_030944	1.57
deoxyribonuclease I-like 3	NM_004944	1.57
ZY6 homolog	BE675849	1.57
tubby like protein 2	NM_003323	1.57
BCL2-like 10 (apoptosis facilitator)	NM_020396	1.57
hypothetical protein FLJ20033	NM_017629	1.57
plexin A1	T16388	1.57
zinc finger protein 44 (KOX 7)	AI758888	1.57
matrix Gla protein	AL523275	1.57
Consensus includes gb:AI346341 /FEA=EST /DB_XREF=gi:4083547 /DB_XREF=est:qp50b09.x1 /CLONE=IMAGE:1926425 /UG=Hs.298998 Homo sapiens, Similar to hypothetical protein FL	AI346341	1.57
dystrobrevin, beta	AI672185	1.57
KIAA0874 protein	X80821	1.57
cAMP response element-binding protein CRE-BPa	NM_004904	1.57
ets variant gene 6 (TEL oncogene)	AF041811	1.57
DKFZP564D206 protein	AI927000	1.57
pp32 related 2	NM_012404	1.57
interferon, alpha 16	NM_002173	1.57
cAMP responsive element binding protein-like 1	NM_004381	1.57
neurobeachin	NM_015678	1.57
aldehyde dehydrogenase 3 family, member B1	NM_000694	1.57
sodium channel, nonvoltage-gated 1 alpha	NM_001038	1.57
heterogeneous nuclear ribonucleoprotein U (scaffold attachment factor A)	AF068846	1.57
synaptotagmin V	NM_003180	1.57

Consensus includes gb:AL109666.1 /DEF=Homo sapiens mRNA full length insert cDNA clone EUROIMAGE 35907. /FEA=mRNA /DB_XREF=gi:5689800 /UG=Hs.7242 Homo sapiens mRNA ful	AL109666	1.57
X17013 B subtilis lys gene for diaminopimelate decarboxylase corresponding to nucleotides 350-1345 of X17013 (-5, -M, -3 represent transcript regions 5 prime, Middle, and 3 prime respe	X17013	1.57
like-glycosyltransferase	BF057493	1.57
Consensus includes gb:BG494940 /FEA=EST /DB_XREF=gi:13456455 /DB_XREF=est:602540961F1 /CLONE=IMAGE:4671854 /UG=Hs.554 Sjogren syndrome antigen A2 (60kD, ribonucleo	BG494940	1.57
TATA element modulatory factor 1	BF593908	1.57
matrix metalloproteinase 28	NM_024302	1.57
potassium channel, subfamily K, member 3 (TASK-1)	NM_002246	1.57
signal transducer and activator of transcription 2, 113kD	NM_005419	1.57
M-phase phosphoprotein 6	NM_005792	1.57
T-box 5	NM_000192	1.57
hypothetical protein FLJ12681	NM_022773	1.57
carbamoyl-phosphate synthetase 1, mitochondrial	W80357	1.57
thymidine kinase 2, mitochondrial	BE895437	1.57
calcitonin receptor	AB022177	1.57
spectrin, beta, non-erythrocytic 5	NM_016642	1.57
succinate dehydrogenase complex, subunit C, integral membrane protein, 15kD	BG110532	1.57
hypothetical protein FLJ13615	AF317887	1.57
group III secreted phospholipase A2	NM_015715	1.57
vasoactive intestinal peptide receptor 1	NM_004624	1.57
follistatin	NM_006350	1.57
myosin VIIA (Usher syndrome 1B (autosomal recessive, severe))	U55209	1.57
hypothetical protein PRO2958	AF119909	1.57
BTB and CNC homology 1, basic leucine zipper transcription factor 1	AF026199	1.57
sex hormone-binding globulin	AI056852	1.57
Rho GTPase activating protein 6	NM_006125	1.57
FGFR1 oncogene partner	NM_007045	1.57
KIAA0356 gene product	AB002354	1.57
DKFZP434N043 protein	AB032967	1.57
catenin (cadherin-associated protein), delta 2 (neural plakophilin-related arm-repeat protein)	U96136	1.57
Consensus includes gb:AV704232 /FEA=EST /DB_XREF=gi:10721550 /DB_XREF=est:AV704232 /CLONE=ADBBLB11 /UG=Hs.228096 Homo sapiens cDNA FLJ11682 fis, clone HEMBA1004E	AV704232	1.57
Consensus includes gb:AK024871.1 /DEF=Homo sapiens cDNA: FLJ21218 fis, clone COL00537. /FEA=mRNA /DB_XREF=gi:10437282 /UG=Hs.306712 Homo sapiens cDNA: FLJ21218 fis, cl	AK024871	1.57
hypothetical protein FLJ20674	NM_019086	1.57
MHC class II transactivator	U18288	1.57
distal-less homeo box 2	BE792224	1.57
chromosome 16 open reading frame 44	NM_024731	1.57
poliovirus receptor	BE615277	1.57
tousled-like kinase 1	AF162666	1.57
urotensin 2	NM_021995	1.57
spondin 1, (f-spondin) extracellular matrix protein	AI885290	1.57
RuvB-like 1 (E. coli)	NM_002784	1.57
hypothetical protein PP5395	NM_021732	1.57
hypothetical protein FLJ10572	NM_018143	1.57
hypothetical protein FLJ22269	AK025922	1.57
Consensus includes gb:N32526 /FEA=EST /DB_XREF=gi:1152925 /DB_XREF=est:yy11f04.s1 /CLONE=IMAGE:270943 /UG=Hs.300642 serologically defined colon cancer antigen 8	N32526	1.57
caudal type homeo box transcription factor 2	U51096	1.57
ankyrin 1, erythrocytic	NM_000037	1.57
neuronal potassium channel alpha subunit	NM_014379	1.57
hypothetical protein PP1345	NM_031245	1.57
H3 histone family, member I	AL522145	1.57
hemoglobin, delta	NM_000519	1.57
a disintegrin and metalloproteinase domain 11	NM_002390	1.57
crystallin, alpha B	AF007162	1.57
gb:BC000772.1 /DEF=Homo sapiens, Similar to KIAA0377 gene product, clone MGC:3268, mRNA, complete cds. /FEA=mRNA /PROD=Similar to KIAA0377 gene product /DB_XREF=gi:12653	BC000772	1.57
immunoglobulin heavy constant mu	AF015124	1.57
gastrulation brain homeo box 1	L11239	1.57
RAB14, member RAS oncogene family	AL162081	1.57
SRY (sex determining region Y)-box 2	AI356682	1.57
KIAA0692 protein	AK025933	1.57
aspartyl-tRNA synthetase	NM_001349	1.57
6 antigen 6	NM_001476	1.57
hypothetical protein FLJ10355	NM_018070	1.57
regulator of G-protein signalling 1	NM_002922	1.57
interferon, alpha-inducible protein (clone IFI-6-16)	NM_022873	1.57

tudor domain containing 1	NM_031278	1.57
NIMA (never in mitosis gene a)-related kinase 1	AF155113	1.57
gb:NM_012348.1 /DEF=Homo sapiens olfactory receptor 89 (OLFR89), mRNA. /FEA=CDS /GEN=OLFR89 /PROD=olfactory receptor 89 /DB_XREF=gi:6912547 /UG=Hs.248225 olfactory re	NM_012348	1.57
Consensus includes gb:AW444520 /FEA=EST /DB_XREF=gi:6986282 /DB_XREF=est:UI-H-BI3-akb-h-11-0-UI.s1 /CLONE=IMAGE:2733933 /UG=Hs.256986 ESTs, Moderately similar to VPP2	AW444520	1.57
granzyme K (serine protease, granzyme 3; tryptase II)	NM_002104	1.57
olfactory receptor, family 51, subfamily E, member 2	NM_030774	1.57
hypothetical protein FLJ10201	NM_018023	1.57
KIAA0789 gene product	NM_014653	1.57
a disintegrin and metalloproteinase domain 7	AF215824	1.57
Consensus includes gb:AA535066 /FEA=EST /DB_XREF=gi:2279319 /DB_XREF=est:nf84a10.s1 /CLONE=IMAGE:926586 /UG=Hs.298553 Homo sapiens, clone IMAGE:3953631, mRNA, part	AA535066	1.57
hypothetical protein DKFZp434D0421	NM_030798	1.57
ectonucleotide pyrophosphatase/phosphodiesterase 2 (autotaxin)	D45421	1.57
small inducible cytokine subfamily C, member 1 (lymphotactin)	NM_002995	1.57
calsequestrin 1 (fast-twitch, skeletal muscle)	NM_001231	1.56
Consensus includes gb:AK024109.1 /DEF=Homo sapiens cDNA FLJ14047 fis, clone HEMBA1006617. /FEA=mRNA /DB_XREF=gi:10436407 /UG=Hs.301186 Homo sapiens cDNA FLJ14047 f	AK024109	1.56
thyrotropin-releasing hormone degrading ectoenzyme	NM_013381	1.56
RAS protein activator like 2	NM_004841	1.56
lymphotoxin alpha (TNF superfamily, member 1)	NM_000595	1.56
Machado-Joseph disease (spinocerebellar ataxia 3, olivopontocerebellar ataxia 3, autosomal dominant, ataxin 3)	NM_004993	1.56
collagen, type VII, alpha 1 (epidermolysis bullosa, dystrophic, dominant and recessive)	NM_000094	1.56
PRO1992 protein	AF118086	1.56
TNF receptor-associated factor 6	NM_004620	1.56
integrin, beta 3 (platelet glycoprotein IIIa, antigen CD61)	U95204	1.56
neuritin	NM_016588	1.56
mitogen-activated protein kinase kinase kinase 8	NM_005204	1.56
hypothetical protein	NM_013386	1.56
Consensus includes gb:AW576457 /FEA=EST /DB_XREF=gi:7247996 /DB_XREF=est:UI-HF-BR0p-aji-a-01-0-UI.s1 /CLONE=IMAGE:3074568 /UG=Hs.20952 Homo sapiens clone 24411 mRN	AW576457	1.56
interleukin 13	NM_002188	1.56
PR domain containing 8	NM_020226	1.56
Consensus includes gb:AL133618.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434C2021 (from clone DKFZp434C2021). /FEA=mRNA /DB_XREF=gi:6599233 /UG=Hs.306443 Homo sapiens	AL133618	1.56
likely ortholog of mouse polydom	NM_024500	1.56
homeo box A3	NM_030661	1.56
gb:NM_003483.2 /DEF=Homo sapiens high-mobility group (nonhistone chromosomal) protein isoform I-C (HMGIC), mRNA. /FEA=mRNA /GEN=HMGIC /PROD=high-mobility group (nonhistc	NM_003483	1.56
Consensus includes gb:AK026847.1 /DEF=Homo sapiens cDNA: FLJ23194 fis, clone REC00490. /FEA=mRNA /DB_XREF=gi:10439802 /UG=Hs.306887 Homo sapiens cDNA: FLJ23194 fis, cl	AK026847	1.56
transketolase-like 1	X91817	1.56
SA (rat hypertension-associated) homolog	D16350	1.56
Rec8p, a meiotic recombination and sister chromatid cohesion phosphoprotein of the rad21p family	NM_005132	1.56
hypothetical protein FLJ12704	NM_024998	1.56
hypothetical protein FLJ23342	NM_024631	1.56
succinate-CoA ligase, ADP-forming, beta subunit	NM_003850	1.56
pregnancy specific beta-1-glycoprotein 4	NM_002780	1.56
collagen, type I, alpha 2	AA788711	1.56
hypothetical protein MGC5601	NM_025247	1.56
KIAA1887 protein	BF446672	1.56
hypothetical protein FLJ11117	NM_018329	1.56
arachidonate 12-lipoxygenase	NM_000697	1.56
hypothetical protein CG003	W58342	1.56
protein phosphatase 1, regulatory (inhibitor) subunit 12B	AF324888	1.56
protein tyrosine phosphatase, receptor type, F	AU145351	1.56
calpain 3, (p94)	BC003169	1.56
discs, large (Drosophila) homolog 3 (neuroendocrine-dlg)	NM_021120	1.56
catenin (cadherin-associated protein), delta 2 (neural plakophilin-related arm-repeat protein)	AF035302	1.56
Consensus includes gb:AU145394 /FEA=EST /DB_XREF=gi:11006915 /DB_XREF=est:AU145394 /CLONE=HEMBA1004711 /UG=Hs.287438 Homo sapiens cDNA FLJ11669 fis, clone HEMBA	AU145394	1.56
IM1 protein	BF224247	1.56
aquaporin 4	U34846	1.56
fasciculation and elongation protein zeta 1 (zygin I)	NM_005103	1.56
cadherin 11, type 2, OB-cadherin (osteoblast)	NM_001797	1.56
Consensus includes gb:AK000220.1 /DEF=Homo sapiens cDNA FLJ20213 fis, clone COLF1971, highly similar to AB007921 Homo sapiens mRNA for KIAA0452 protein. /FEA=mRNA /DB_XRE	AK000220	1.56
phosphodiesterase 7B	NM_018945	1.56
hypothetical protein FLJ20087	NM_017662	1.56
glycoprotein M6B	AF016004	1.56
discs, large (Drosophila) homolog 5	BC002915	1.56
Consensus includes gb:N71739 /FEA=EST /DB_XREF=gi:1228451 /DB_XREF=est:yw52h11.r1 /CLONE=IMAGE:255909 /UG=Hs.237554 Homo sapiens cDNA FLJ14306 fis, clone PLACE3000	N71739	1.56

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v-erb-b2 erythroblastic leukemia viral oncogene homolog 3 (avian)	NM_001982	1.56
Consensus includes gb:AV700403 /FEA=EST /DB_XREF=gi:10302374 /DB_XREF=est:AV700403 /CLONE=GKBADA01 /UG=Hs.293673 ESTs	AV700403	1.56
amelogenin (Y chromosome)	NM_001143	1.56
Consensus includes gb:BE673151 /FEA=EST /DB_XREF=gi:10033692 /DB_XREF=est:7d30a11.x1 /CLONE=IMAGE:3248732 /UG=Hs.91626 Homo sapiens clone 818 unknown mRNA	BE673151	1.56
tumor endothelial marker 5 precursor	BF511315	1.56
homeo box 11 (T-cell lymphoma 3-associated breakpoint)	NM_005521	1.56
prostate tumor over expressed gene 1	U79287	1.56
opioid receptor, mu 1	NM_000914	1.56
glutamate decarboxylase 1 (brain, 67kD)	NM_013445	1.56
ubiquitin-like 1 (sentrin)	BC006462	1.56
eukaryotic translation initiation factor 4E	NM_001968	1.56
Hin-1	NM_017493	1.56
apolipoprotein L, 5	NM_030642	1.56
Consensus includes gb:AK025323.1 /DEF=Homo sapiens cDNA: FLJ21670 fis, clone COL09010. /FEA=mRNA /DB_XREF=gi:10437816 /UG=Hs.230999 Homo sapiens cDNA: FLJ21670 fis, cl	AK025323	1.56
flavin containing monooxygenase 3	NM_006894	1.56
homeo box A9	AI246769	1.56
contactin associated protein-like 2	AB020675	1.56
bone morphogenetic protein 8 (osteogenic protein 2)	NM_001720	1.56
KIAA0300 protein	AF338650	1.56
F-box only protein 22	NM_012170	1.56
bromodomain-containing 3	NM_007371	1.56
WAS protein family, member 3	AB020707	1.56
lymphocyte antigen 117	NM_007161	1.56
paired related homeobox protein	NM_016307	1.56
hypothetical protein PRO2958	NM_018546	1.56
6 protein-coupled receptor 49	AF062006	1.56
KIAA1079 protein	NM_014916	1.56
human epididymis-specific 3 beta	NM_022360	1.56
membrane-spanning 4-domains, subfamily A, member 12 4-domains, subfamily A, member 7	NM_017716	1.56
splicing factor 3a, subunit 2, 66kD	L21990	1.56
Consensus includes gb:AL049260.1 /DEF=Homo sapiens mRNA; cDNA DKFZp564E233 (from clone DKFZp564E233). /FEA=mRNA /DB_XREF=gi:4500007 /UG=Hs.302050 Homo sapiens mF	AL049260	1.56
lipophilin B (uteroglobin family member), prostatein-like	NM_006551	1.56
DKFZP586M1019 protein	NM_015603	1.56
acetylserotonin O-methyltransferase	BC001620	1.56
chromosome X open reading frame 6	NM_005491	1.56
high-mobility group 20B	BC002552	1.56
nuclear receptor coactivator 1	U19179	1.56
serum constituent protein	NM_007061	1.56
collagen, type IV, alpha 6	AI889941	1.56
I factor (complement)	NM_000204	1.56
Consensus includes gb:AA888858 /FEA=EST /DB_XREF=gi:3015728 /DB_XREF=est:ak23d05.s1 /CLONE=IMAGE:1406793 /UG=Hs.118567 ESTs	AA888858	1.56
KIAA0087 gene product	NM_014769	1.56
preferentially expressed antigen in melanoma	NM_006115	1.55
hypothetical protein MGC13053	BC006134	1.55
Consensus includes gb:X69637.1 /DEF=H.sapiens mRNA sequence (16p11.2). /FEA=mRNA /DB_XREF=gi:29573 /UG=Hs.307208 H.sapiens mRNA sequence (16p11.2)	X69637	1.55
folate hydrolase (prostate-specific membrane antigen) 1	NM_004476	1.55
leukocyte membrane antigen	AF020314	1.55
glucose phosphate isomerase	AF187554	1.55
hypothetical protein PRO2130	NM_018513	1.55
POU domain, class 6, transcription factor 1	NM_002702	1.55
double ring-finger protein, Dorfin	NM_015435	1.55
CED-6 protein	AK023668	1.55
thyroid stimulating hormone, beta	NM_000549	1.55
Consensus includes gb:AL117643.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434M245 (from clone DKFZp434M245). /FEA=mRNA /DB_XREF=gi:5912233 /UG=Hs.5288 Homo sapiens mRN	AL117643	1.55
Krueppel-related zinc finger protein	NM_015852	1.55
hypothetical protein MGC5528	L11372	1.55
tomosyn	AB023223	1.55
Consensus includes gb:AA365670 /FEA=EST /DB_XREF=gi:2018010 /DB_XREF=est:EST76509 /UG=Hs.261526 ESTs	AA365670	1.55
Consensus includes gb:AK024871.1 /DEF=Homo sapiens cDNA: FLJ21218 fis, clone COL00537. /FEA=mRNA /DB_XREF=gi:10437282 /UG=Hs.306712 Homo sapiens cDNA: FLJ21218 fis, cl	AK024871	1.55
a disintegrin and metalloproteinase domain 22	NM_004194	1.55
insulin-like growth factor 1 (somatomedin C)	AU144912	1.55
fucosyltransferase 4 (alpha (1,3) fucosyltransferase, myeloid-specific)	AF305083	1.55
DKFZP586N0819 protein	NM_015437	1.55

ras-related C3 botulinum toxin substrate 1 (rho family, small GTP binding protein Rac1)	BC004247	1.55
platelet/endothelial cell adhesion molecule (CD31 antigen)	AW574504	1.55
melanoma antigen, family D, 2	AI924630	1.55
hepatocyte growth factor (hepapoietin A; scatter factor)	M77227	1.55
insulin-like growth factor binding protein 3	M31159	1.55
fibroblast growth factor 20	NM_019851	1.55
low density lipoprotein-related protein 2	NM_004525	1.55
epsin 3	NM_017957	1.55
immunoglobulin superfamily, member 6	NM_005849	1.55
HCF-binding transcription factor Zhangfei	AA521272	1.55
transmembrane, prostate androgen induced RNA	NM_020182	1.55
cyclin-dependent kinase-like 2 (CDC2-related kinase)	NM_003948	1.55
nuclear receptor co-repressor 1	BF437948	1.55
glutamate-ammonia ligase (glutamine synthase)	NM_002065	1.55
glucagon-like peptide 2 receptor	NM_004246	1.55
poly(A) polymerase gamma	NM_022894	1.55
Consensus includes gb:AL359052.1 /DEF=Homo sapiens mRNA full length insert cDNA clone EUROIMAGE 1968422. /FEA=mRNA /PROD=ITGBL1, integrin beta-like 1 /DB_XREF=gi:8518175	AL359052	1.55
hypothetical protein FLJ14033 similar to hypoxia inducible factor 3, alpha subunit	AK021881	1.55
DEAD/H (Asp-Glu-Ala-Asp/His) box binding protein 1	AJ348378	1.55
insulin	NM_000207	1.55
regulator of G-protein signalling 12	AF030111	1.55
hypothetical protein FLJ12891	NM_024950	1.55
heat shock 27kD protein 2	NM_001541	1.55
E74-like factor 5 (ets domain transcription factor)	NM_001422	1.55
KIAA0480 gene product	NM_014810	1.55
Fc fragment of IgG, high affinity Ia, receptor for (CD64)	X14355	1.55
KIAA1005 protein	BF515597	1.55
hypothetical protein FLJ12987	NM_025170	1.55
macrophage scavenger receptor 1	AI299239	1.55
heterogeneous nuclear ribonucleoprotein H1 (H)	AI022387	1.55
SPR1 protein	NM_014069	1.55
bone morphogenetic protein 1	NM_006129	1.55
prostaglandin E receptor 3 (subtype EP3)	NM_000957	1.55
hemochromatosis	AF144243	1.55
trinucleotide repeat containing 9	AK025084	1.55
transcription elongation factor A (SII), 2	NM_003195	1.55
microtubule-associated protein 1A	AF095910	1.55
gap junction protein, beta 3, 31kD (connexin 31)	NM_024009	1.55
Down syndrome critical region gene 4	NM_005867	1.55
hypothetical protein PRO1995	NM_018612	1.55
ATP-binding cassette, sub-family C (CFTR/MRP), member 6	AI074459	1.55
PR domain containing 10	AV682503	1.55
Consensus includes gb:AK021741.1 /DEF=Homo sapiens cDNA FLJ11679 fis, clone HEMBA1004807. /FEA=mRNA /DB_XREF=gi:10432986 /UG=Hs.287439 Homo sapiens cDNA FLJ11679 fi	AK021741	1.55
paraneoplastic antigen	AI422335	1.55
fatty acid binding protein 7, brain	NM_001446	1.55
solute carrier family 22 (organic cation transporter), member 2	NM_003058	1.55
cysteine-rich, angiogenic inducer, 61	NM_001554	1.55
steroidogenic acute regulatory protein	NM_000349	1.55
adrenergic, beta-3-, receptor	NM_000025	1.55
acid phosphatase 5, tartrate resistant	NM_001611	1.55
hairy homolog (Drosophila)	BE973687	1.55
DKFZP434K091 protein	AB031547	1.55
transcription factor AP-4 (activating enhancer binding protein 4)	NM_003223	1.55
tumor necrosis factor receptor superfamily, member 6	AA164751	1.55
retinoschisis (X-linked, juvenile) 1	NM_000330	1.55
NAD(P)H dehydrogenase, quinone 1	AI039874	1.55
HSKM-B protein	AF070592	1.55
variable charge, Y chromosome, 2	NM_004678	1.55
hypothetical protein FLJ10620	NM_018157	1.55
tumor necrosis factor receptor superfamily, member 10c, decoy without an intracellular domain	NM_003841	1.55
FBJ murine osteosarcoma viral oncogene homolog B	NM_006732	1.55
glycine-N-acyltransferase	AW024233	1.55
Consensus includes gb:AU147620 /FEA=EST /DB_XREF=gi:11009141 /DB_XREF=est:AU147620 /CLONE=MAMMA1001206 /UG=Hs.288490 Homo sapiens cDNA FLJ12232 fis, clone MAMM	AU147620	1.55

Consensus includes gb:AK026273.1 /DEF=Homo sapiens cDNA: FLJ22620 fis, clone HSI05629. /FEA=mRNA /DB_XREF=gi:10439077 /UG=Hs.306834 Homo sapiens cDNA: FLJ22620 fis, cl	AK026273	1.55
arachidonate 15-lipoxygenase, second type	NM_001141	1.55
KIAA1466 protein	AJ765383	1.55
hypothetical protein FLJ11506	AL136715	1.55
sulfotransferase family, cytosolic, 1C, member 1	AF026303	1.55
hypothetical protein MGC10848	NM_030569	1.55
SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily a, member 2	X72889	1.55
melanoma antigen, family A, 10	BC004105	1.55
heterogeneous nuclear ribonucleoprotein L	AL044078	1.55
translin	AJ659180	1.55
Bacillus subtilis /REF=L38424 /DEF=B subtilis dapB, jojF, jojG genes corresponding to nucleotides 1439-1846 of L38424 /LEN=1931 (-5, -M, -3 represent transcript regions 5 prime, Middl	L38424	1.55
Consensus includes gb:AF070625.1 /DEF=Homo sapiens clone 24734 mRNA sequence. /FEA=mRNA /DB_XREF=gi:3283891 /UG=Hs.12440 Homo sapiens clone 24734 mRNA sequence	AF070625	1.55
KIAA0590 gene product	NM_014714	1.55
Consensus includes gb:AW974915 /FEA=EST /DB_XREF=gi:8166118 /DB_XREF=est:EST387020 /UG=Hs.116550 ESTs	AW974915	1.55
phosphodiesterase 4B, cAMP-specific (phosphodiesterase E4 dunce homolog, Drosophila)	NM_002600	1.55
zinc finger protein 259, pseudogene	Z95118	1.55
mucin 5, subtype B, tracheobronchial	AJ521646	1.55
hypothetical protein FLJ11196	NM_018357	1.55
LIM domain only 1 (rhombotin 1)	NM_002315	1.55
aquaporin 1 (channel-forming integral protein, 28kD)	NM_000385	1.55
cell recognition molecule CASPR3	NM_024879	1.55
small muscle protein, X-linked	NM_014332	1.55
tumor necrosis factor (ligand) superfamily, member 13	AF114011	1.55
Consensus includes gb:AA707199 /FEA=EST /DB_XREF=gi:2717117 /DB_XREF=est:zj32g01.s1 /CLONE=IMAGE:452016 /UG=Hs.298998 Homo sapiens, Similar to hypothetical protein FLJ2	AA707199	1.55
ATP-binding cassette, sub-family G (WHITE), member 4	NM_022169	1.55
Consensus includes gb:AL035555 /DEF=Human DNA sequence from clone 217P22 on chromosome 6p21.1-21.31. Contains part of a novel Dynein heavy chain family protein gene and a pse	AL035555	1.55
C-type (calcium dependent, carbohydrate-recognition domain) lectin, superfamily member 9	NM_014358	1.55
solute carrier family 22 (organic anion transporter), member 7	NM_006672	1.55
cadherin-like protein VR20	AL109928	1.55
interleukin 1 receptor antagonist	AW083357	1.55
cytochrome P450, subfamily I (aromatic compound-inducible), polypeptide 1	NM_000499	1.55
fatty acid hydroxylase	NM_024306	1.55
mitogen-activated protein kinase kinase 6	NM_002758	1.55
protein phosphatase 3 (formerly 2B), catalytic subunit, gamma isoform (calcineurin A gamma)	AJ762547	1.55
Consensus includes gb:AL134612 /FEA=EST /DB_XREF=gi:6602799 /DB_XREF=est:DKFZp547E086_s1 /CLONE=DKFZp547E086 /UG=Hs.167036 Homo sapiens clone 23798 and 23825 m	AL134612	1.55
N-acetylated alpha-linked acidic dipeptidase 2	AJ012370	1.54
Consensus includes gb:W86781 /FEA=EST /DB_XREF=gi:1400509 /DB_XREF=est:zh64a03.s1 /CLONE=IMAGE:416812 /UG=Hs.293736 ESTs	W86781	1.54
solute carrier family 18 (vesicular acetylcholine), member 3	NM_003055	1.54
discs, large (Drosophila) homolog 2 (chapsyn-110)	NM_001364	1.54
T-cell acute lymphocytic leukemia 1	NM_003189	1.54
protein kinase, X-linked	NM_005044	1.54
PL6 protein	BF034906	1.54
type 1 tumor necrosis factor receptor shedding aminopeptidase regulator	AA195244	1.54
kelch-like 4 (Drosophila)	BF215673	1.54
polycythemia rubra vera 1; cell surface receptor	NM_020406	1.54
killer cell lectin-like receptor subfamily D, member 1	AB009597	1.54
early growth response 4	NM_001965	1.54
ubiquitin specific protease 6 (Tre-2 oncogene)	NM_004505	1.54
early lymphoid activation protein	L22650	1.54
adrenergic, alpha-1D-, receptor	M76446	1.54
nescient helix loop helix 2	NM_005599	1.54
sialic acid binding Ig-like lectin 7	AJ130713	1.54
hypothetical protein DKFZp761P1010	AF251059	1.54
hypothetical protein FLJ11726	NM_024971	1.54
general transcription factor IIH, polypeptide 1 (62kD subunit)	BC000365	1.54
hemochromatosis	AF150664	1.54
hypothetical protein FLJ22615	NM_024919	1.54
cholinergic receptor, nicotinic, beta polypeptide 3	X67513	1.54
RNA binding motif protein 4	BE222257	1.54
hypothetical protein FLJ10038	NM_017976	1.54
Consensus includes gb:AI088162 /FEA=EST /DB_XREF=gi:3427138 /DB_XREF=est:oz96b09.s1 /CLONE=IMAGE:1683161 /UG=Hs.262292 ESTs, Moderately similar to ALU3_HUMAN ALU SI	AI088162	1.54
Consensus includes gb:N34842 /FEA=EST /DB_XREF=gi:1155984 /DB_XREF=est:yy45d11.s1 /CLONE=IMAGE:276501 /UG=Hs.6580 Homo sapiens cDNA: FLJ23227 fis, clone CAE00645, h	N34842	1.54
Consensus includes gb:AK021614.1 /DEF=Homo sapiens cDNA FLJ11552 fis, clone HEMBA1003021. /FEA=mRNA /DB_XREF=gi:10432830 /UG=Hs.306606 Homo sapiens cDNA FLJ11552 fi	AK021614	1.54

transcription factor AP-2 gamma (activating enhancer binding protein 2 gamma)	NM_003222	1.54
chromosome 21 open reading frame 80	NM_015227	1.54
zinc finger protein 236	AA004757	1.54
likely homolog of rat kinase D-interacting substance of 220 kDa	AL133620	1.54
89 protein	BC002944	1.54
osteoglycin (osteoinductive factor, mimecan)	NM_014057	1.54
Consensus includes gb:AI054381 /FEA=EST /DB_XREF=gi:3322168 /DB_XREF=est:qi64d09.x1 /CLONE=IMAGE:1861265 /UG=Hs.176139 ESTs	AI054381	1.54
epithelial membrane protein 1	NM_001423	1.54
fibroblast growth factor receptor 1 (fms-related tyrosine kinase 2, Pfeiffer syndrome)	M63889	1.54
killer cell immunoglobulin-like receptor, three domains, long cytoplasmic tail, 2	NM_006737	1.54
hypothetical protein FLJ14082	NM_025024	1.54
gb:L14457.1 /DEF=Human Ig rearranged kappa-chain gene V-J-region, complete cds. /FEA=mRNA /DB_XREF=gi:398491 /FL=gb:L14457.1	L14457	1.54
hypothetical protein FLJ12331	NM_024986	1.54
Vertebrate LIN7 homolog 1, Tax interaction protein 33	NM_004664	1.54
hypothetical protein MGC2652	NM_024328	1.54
transcriptional co-activator with PDZ-binding motif (TAZ)	AA081084	1.54
KIAA0903 protein	BF116032	1.54
phosphatidylinositol glycan, class K	AF022913	1.54
claudin 8	AL049977	1.54
Consensus includes gb:AF164963.1 /DEF=Homo sapiens tumor antigen NA88-A pseudogene, complete sequence. /FEA=mRNA /DB_XREF=gi:5901726 /UG=Hs.306576 Homo sapiens tumo	AF164963	1.54
Consensus includes gb:AK023515.1 /DEF=Homo sapiens cDNA FLJ13453 fis, clone PLACE1003205. /FEA=mRNA /DB_XREF=gi:10435471 /UG=Hs.301715 Homo sapiens cDNA FLJ13453 fi	AK023515	1.54
hypothetical protein FLJ14033 similar to hypoxia inducible factor 3, alpha subunit	NM_022462	1.54
geranylgeranyl diphosphate synthase 1	AW299507	1.54
double-stranded RNA specific adenosine deaminase	NM_018702	1.54
KIAA1025 protein	AL133033	1.54
selectin E (endothelial adhesion molecule 1)	NM_000450	1.54
glutamate receptor, metabotropic 7	NM_000844	1.54
similar to rat tricarboxylate carrier-like protein	NM_030971	1.54
CD3-epsilon-associated protein; antisense to ERCC-1	NM_012099	1.54
UDP-glucose dehydrogenase	NM_003359	1.54
zinc finger protein 37a (KOX 21)	AU118165	1.54
postmeiotic segregation increased 2-like 3	D38437	1.54
zinc finger protein 268	AF317549	1.54
testis specific protein, Y-linked	M94893	1.54
guanine nucleotide binding protein (G protein), alpha activating activity polypeptide O	AA758795	1.54
protein tyrosine phosphatase, receptor type, H	NM_002842	1.54
sodium channel, voltage-gated, type X, alpha polypeptide	NM_006514	1.54
protease, serine, 16 (thymus)	NM_005865	1.54
ring finger protein 3	NM_006315	1.54
tripartite motif-containing 2	NM_015271	1.54
Consensus includes gb:AC006986 /DEF=Homo sapiens BAC clone RP11-155I5 from Y /FEA=CDS /DB_XREF=gi:4753246 /UG=Hs.283908 Homo sapiens BAC clone RP11-155I5 from Y	AC006986	1.54
acid fibroblast growth factor-like protein	NM_013394	1.54
adenomatous polyposis coli like	NM_005883	1.54
hypothetical protein PRO2435	NM_018527	1.54
small inducible cytokine B subfamily (Cys-X-Cys motif), member 13 (B-cell chemoattractant)	NM_006419	1.54
N-myristoyltransferase 1	NM_021079	1.54
thymine-DNA glycosylase	BF674842	1.54
thymidylate synthetase	BG281679	1.54
tissue factor pathway inhibitor (lipoprotein-associated coagulation inhibitor)	AF021834	1.54
Consensus includes gb:AV700298 /FEA=EST /DB_XREF=gi:10302269 /DB_XREF=est:AV700298 /CLONE=GKCBVG05 /UG=Hs.285173 ESTs, Moderately similar to CD44_HUMAN CD44 ANTIG	AV700298	1.54
chloride channel, calcium activated, family member 2	NM_006536	1.54
FAT tumor suppressor homolog 1 (Drosophila)	NM_005245	1.54
Consensus includes gb:AA045174 /FEA=EST /DB_XREF=gi:1523376 /DB_XREF=est:zk66a07.s1 /CLONE=IMAGE:487764 /UG=Hs.16193 Homo sapiens mRNA; cDNA DKFZp586B211 (from	AA045174	1.54
KIAA1026 protein	AB028949	1.54
MAD, mothers against decapentaplegic homolog 9 (Drosophila)	NM_005905	1.54
hypothetical protein	NM_018985	1.54
SEEK1 protein	NM_014068	1.54
hypothetical protein FLJ14280	NM_024886	1.54
caspase 1, apoptosis-related cysteine protease (interleukin 1, beta, convertase)	M87507	1.54
dynamin 1	L07810	1.54
adenylate kinase 5	NM_012093	1.54
Hermansky-Pudlak syndrome	NM_000195	1.54
glucagon-like peptide 1 receptor	U01157	1.54

type 1 tumor necrosis factor receptor shedding aminopeptidase regulator	AF183569	1.54
voltage-gated sodium channel beta-3 subunit (scn3b gene)	AW007335	1.54
6 protein-coupled receptor	U73531	1.54
Consensus includes gb:AW156899 /FEA=EST /DB_XREF=gi:6228300 /DB_XREF=est:au89c11.x1 /CLONE=IMAGE:2783444 /UG=Hs.12520 Homo sapiens clone 23568, 23621, 23795, 238	AW156899	1.54
laminin, beta 4	AF029325	1.54
hypothetical protein FLJ22087	NM_022070	1.54
chromosome 4 open reading frame 5	BF939176	1.54
Consensus includes gb:AA621286 /FEA=EST /DB_XREF=gi:2525225 /DB_XREF=est:zu83e09.s1 /CLONE=IMAGE:744616 /UG=Hs.179697 Human DNA damage repair and recombination pro	AA621286	1.54
arginine vasopressin (neurophysin II, antidiuretic hormone, diabetes insipidus, neurohypophyseal)	NM_000490	1.54
alpha-catenin-like protein	NM_013266	1.54
hypothetical protein FLJ12969	NM_022838	1.54
purinergic receptor P2Y, G-protein coupled, 2	NM_002564	1.54
RAB6 interacting, kinesin-like (rabkinesin 6)	NM_005733	1.54
hypothetical protein	NM_019016	1.54
FERM, RhoGEF (ARHGEF) and pleckstrin domain protein 1 (chondrocyte-derived)	NM_005766	1.54
hypothetical protein FLJ13511	AF209931	1.54
genethonin 1	NM_003943	1.54
multiple inositol polyphosphate histidine phosphatase, 1	AF084943	1.54
interleukin 1, epsilon	NM_014440	1.54
PDZ and LIM domain 1 (elfin)	BC000915	1.53
son of sevenless (Drosophilia) homolog 2	L20686	1.53
Cluster Incl. AI400621:tg49g02.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-2112146 /clone_end=3 /gb=AI400621 /gi=4243708 /ug=Hs.213640 /len=529	AI400621	1.53
hypothetical protein FLJ11259	NM_018370	1.53
Consensus includes gb:S83374.1 /DEF=glutamate transporter II variant BHBGT IIB (5 region) human, brain and spinal cord, mRNA Partial Mutant, 129 nt. /FEA=mRNA /DB_XREF=gi:191162	S83374	1.53
solute carrier family 8 (sodium/calcium exchanger), member 1	AF108389	1.53
5-hydroxytryptamine (serotonin) receptor 4	AI131724	1.53
Consensus includes gb:AK022331.1 /DEF=Homo sapiens cDNA FLJ12269 fis, clone MAMMA1001635. /FEA=mRNA /DB_XREF=gi:10433706 /UG=Hs.293925 Homo sapiens cDNA FLJ12269	AK022331	1.53
KIAA0902 protein	NM_014927	1.53
oxytocin receptor	NM_000916	1.53
glycoprotein M6B	AW148844	1.53
hypothetical protein FLJ14251	NM_024881	1.53
sphingomyelin phosphodiesterase 2, neutral membrane (neutral sphingomyelinase)	AI739480	1.53
PHD finger protein 2	NM_005392	1.53
glutamate receptor, ionotropic, kainate 3	NM_000831	1.53
Consensus includes gb:U05589.1 /DEF=Human ribosomal protein S1 homolog mRNA, partial cds. /FEA=mRNA /DB_XREF=gi:497001 /UG=Hs.371 ribosomal protein S1-like	U05589	1.53
DNAJ domain-containing	NM_013238	1.53
extracellular glycoprotein EMILIN-2 precursor	AL117592	1.53
SFRS protein kinase 2	AW149364	1.53
immunoglobulin heavy constant gamma 3 (G3m marker)	AJ275408	1.53
keratin 1 (epidermolytic hyperkeratosis)	NM_006121	1.53
Kruppel-like factor 12	NM_016285	1.53
nectin 3	AA129716	1.53
protein tyrosine phosphatase, receptor type, T	NM_007050	1.53
fibroblast growth factor 1 (acidic)	X59065	1.53
zinc finger protein 177	NM_003451	1.53
tetraspan NET-6 protein	NM_014399	1.53
lectin, galactoside-binding, soluble, 8 (galectin 8)	AF074000	1.53
chromosome 17 open reading frame 1A	U43383	1.53
Zic family member 1 (odd-paired homolog, Drosophila)	NM_003412	1.53
tubby like protein 3	AK024246	1.53
stanniocalcin 1	NM_003155	1.53
chloride channel, calcium activated, family member 2	BF003134	1.53
kallikrein 7 (chymotryptic, stratum corneum)	NM_005046	1.53
WD repeat domain 4	BC001074	1.53
anterior gradient 2 (Xenopus laevis) homolog	AF088867	1.53
hypothetical protein MGC2615	NM_024103	1.53
Consensus includes gb:AW772192 /FEA=EST /DB_XREF=gi:7704256 /DB_XREF=est:hn69f02.x1 /CLONE=IMAGE:3033147 /UG=Hs.7888 Homo sapiens clone 23736 mRNA sequence	AW772192	1.53
retinoblastoma binding protein 6	NM_006910	1.53
olfactory receptor, family 7, subfamily E, member 16 pseudogene	AC006271	1.53
cardiac ankyrin repeat protein	NM_014391	1.53
Consensus includes gb:AJ243936.1 /DEF=Homo sapiens mRNA for G16 protein (G16 gene located in the class III region of the major histocompatibility complex). /FEA=mRNA /GEN=G16 /P	AJ243936	1.53
Consensus includes gb:AK024606.1 /DEF=Homo sapiens cDNA: FLJ20953 fis, clone ADSE01979. /FEA=mRNA /DB_XREF=gi:10436920 /UG=Hs.306697 Homo sapiens cDNA: FLJ20953 fis,	AK024606	1.53
S164 protein	BF055107	1.53

DKFZP564A122 protein	AL110243	1.53
RAB6B, member RAS oncogene family	AW118072	1.53
solute carrier family 26 (sulfate transporter), member 2	AI025519	1.53
leptin (murine obesity homolog)	NM_000230	1.53
Consensus includes gb:AL080180.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434M091 (from clone DKFZp434M091). /FEA=mRNA /DB_XREF=gi:5262656 /UG=Hs.182293 Homo sapiens m	AL080180	1.53
CD84 antigen (leukocyte antigen)	AF054816	1.53
death-associated protein kinase 2	NM_014326	1.53
Consensus includes gb:AW851559 /FEA=EST /DB_XREF=gi:7947076 /DB_XREF=est:IL3-CT0221-040400-107-E09 /UG=Hs.202695 Human soluble CD44 (CD44) mRNA, with exon v9 extens	AW851559	1.53
four and a half LIM domains 1	U29538	1.53
cancer/testis antigen	U87459	1.53
kallikrein 13	AL050220	1.53
mannosyl (alpha-1,6-)-glycoprotein beta-1,2-N-acetylglucosaminyltransferase	NM_002408	1.53
Leman coiled-coil protein	NM_016201	1.53
keratin associated protein 1.1	NM_030967	1.53
tumor necrosis factor (ligand) superfamily, member 12	NM_003809	1.53
hypothetical protein MGC10818	NM_030568	1.53
UDP-Gal:betaGlcNAc beta 1,3-galactosyltransferase, polypeptide 1	NM_020981	1.53
lipoma HMGIC fusion partner	NM_005780	1.53
ADP-ribosylation factor 4-like	NM_001661	1.53
claudin 4	NM_001305	1.53
fibroblast growth factor 5	NM_004464	1.53
Consensus includes gb:AL049992.1 /DEF=Homo sapiens mRNA; cDNA DKFZp564G242 (from clone DKFZp564G242). /FEA=mRNA /DB_XREF=gi:4884244 /UG=Hs.225836 Homo sapiens mI	AL049992	1.53
actin like protein	BF594459	1.53
melanoma antigen, family A, 4	AW438674	1.53
a disintegrin and metalloproteinase domain 12 (meltrin alpha)	NM_003474	1.53
vanin 3	NM_018399	1.53
glucagon-like peptide 1 receptor	NM_002062	1.53
Consensus includes gb:L37198.1 /DEF=Homo sapiens (clone B3B3E13) Huntingtons disease candidate region mRNA fragment. /FEA=mRNA /DB_XREF=gi:600519 /UG=Hs.233617 Homo s	L37198	1.53
transforming growth factor, alpha	AF149096	1.53
Consensus includes gb:N54942 /FEA=EST /DB_XREF=gi:1196262 /DB_XREF=est:yv38b08.s1 /CLONE=IMAGE:244983 /UG=Hs.276590 ESTs	N54942	1.53
homeo box C6	NM_004503	1.53
dead ringer (Drosophila)-like 2 (bright and dead ringer)	NM_006465	1.53
hypothetical protein FLJ11827	NM_025093	1.53
serine/arginine repetitive matrix 2	AI655799	1.53
hypothetical protein FLJ11136	AI497781	1.53
small inducible cytokine A5 (RANTES)	NM_002985	1.53
uncharacterized bone marrow protein BM040	NM_018456	1.53
immunoglobulin superfamily, member 4	AF132811	1.53
membrane-spanning 4-domains, subfamily A, member 5	NM_023945	1.53
dickkopf homolog 1 (Xenopus laevis)	NM_012242	1.53
prolactin	NM_000948	1.53
Sjogren syndrome antigen A2 (60kD, ribonucleoprotein autoantigen SS-A/Ro)	AK024044	1.53
hypothetical protein FLJ20378	NM_017795	1.53
DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide, Y chromosome	AF000985	1.53
fibroblast growth factor 16	NM_003868	1.53
brain-specific angiogenesis inhibitor 3	NM_001704	1.53
sema domain, immunoglobulin domain (Ig), short basic domain, secreted, (semaphorin) 3C	AJ962897	1.53
NK6 transcription factor homolog A (Drosophila)	NM_006168	1.53
phytanoyl-CoA hydroxylase interacting protein	NM_014759	1.53
fucosyltransferase 4 (alpha (1,3) fucosyltransferase, myeloid-specific)	M58596	1.53
CD3Z antigen, zeta polypeptide (TiT3 complex)	J04132	1.53
transcriptional activator of the c-fos promoter	NM_006365	1.53
hypothetical protein FLJ20255	NM_017728	1.53
hypothetical protein FLJ10134	NM_018004	1.53
Consensus includes gb:N21364 /FEA=EST /DB_XREF=gi:1126534 /DB_XREF=est:yz50h10.s1 /CLONE=IMAGE:265219 /UG=Hs.26968 Homo sapiens mRNA from chromosome 5q21-22, clo	N21364	1.53
testis specific protein, Y-linked	NM_003308	1.53
nucleobindin 2	NM_005013	1.53
hypothetical protein FLJ13265	NM_024877	1.53
HT017 protein	AF225421	1.53
potassium inwardly-rectifying channel, subfamily J, member 12	NM_021012	1.53
CCR4-NOT transcription complex, subunit 4	U71268	1.53
C-type (calcium dependent, carbohydrate-recognition domain) lectin, superfamily member 12	AF313468	1.53
mannosyl (alpha-1,3-)-glycoprotein beta-1,4-N-acetylglucosaminyltransferase, isoenzyme A	NM_012214	1.53

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metastasis-associated 1-like 1	NM_004739	1.53
tryptophan 2,3-dioxygenase	NM_005651	1.53
phosphorylase kinase, alpha 1 (muscle)	NM_002637	1.53
unc-5 homolog B (C. elegans)	NM_003728	1.53
Consensus includes gb:AL390149.1 /DEF=Homo sapiens mRNA; cDNA DKFZp5478026 (from clone DKFZp5478026). /FEA=mRNA /DB_XREF=gi:9368883 /UG=Hs.283862 Homo sapiens mI	AL390149	1.53
tripartite motif-containing 14	BC006333	1.53
zinc finger protein 74 (Cos52)	NM_003426	1.53
Consensus includes gb:AU148255 /FEA=EST /DB_XREF=gi:11009776 /DB_XREF=est:AU148255 /CLONE=MAMMA1003031 /UG=Hs.287627 Homo sapiens cDNA FLJ14151 fis, clone MAMM	AU148255	1.53
Consensus includes gb:AK023345.1 /DEF=Homo sapiens cDNA FLJ13283 fis, clone OVARC1001113, highly similar to Homo sapiens diaphanous 1 (HDIA1) mRNA. /FEA=mRNA /DB_XREF=g	AK023345	1.53
vasoactive intestinal peptide receptor 2	NM_003382	1.53
T cell receptor delta locus	X06557	1.53
protein kinase, Y-linked	NM_002760	1.53
cytochrome P450, subfamily IIC (mephenytoin 4-hydroxylase), polypeptide 18	NM_000772	1.53
integrin, beta 5	AL048423	1.53
KIAA0450 gene product	NM_014638	1.53
hypothetical protein FLJ22059	NM_022752	1.53
Consensus includes gb:AK022363.1 /DEF=Homo sapiens cDNA FLJ12301 fis, clone MAMMA1001858. /FEA=mRNA /DB_XREF=gi:10433745 /UG=Hs.287511 Homo sapiens cDNA FLJ12301 f	AK022363	1.53
c-mer proto-oncogene tyrosine kinase	NM_006343	1.53
KIAA0716 gene product	NM_014705	1.53
melanoma antigen, family A, 1 (directs expression of antigen MZ2-E)	NM_004988	1.52
putative tumor suppressor	BC004483	1.52
Consensus includes gb:AV763520 /FEA=EST /DB_XREF=gi:10921368 /DB_XREF=est:AV763520 /CLONE=MDSCIH03 /UG=Hs.282113 ESTs	AV763520	1.52
dom-3 homolog 2 (C. elegans)	AF059252	1.52
likely ortholog of mouse piwi like homolog 1 (Drosophila)-like	NM_018068	1.52
matrix metalloproteinase 8 (neutrophil collagenase)	NM_002424	1.52
Consensus includes gb:AK025152.1 /DEF=Homo sapiens cDNA: FLJ21499 fis, clone COL05634. /FEA=mRNA /DB_XREF=gi:10437612 /UG=Hs.306773 Homo sapiens cDNA: FLJ21499 fis, cl	AK025152	1.52
phorbolin-like protein MDS019	AK022802	1.52
hypothetical protein FLJ11042	NM_018308	1.52
receptor-interacting serine-threonine kinase 2	AF027706	1.52
Consensus includes gb:BE910600 /FEA=EST /DB_XREF=gi:10407359 /DB_XREF=est:601501171F1 /CLONE=IMAGE:3903161 /UG=Hs.288751 ESTs, Highly similar to HIC protein isoform p	BE910600	1.52
vitamin D (1,25- dihydroxyvitamin D3) receptor	AA904259	1.52
calpain 2, (m/II) large subunit	M23254	1.52
hypothetical protein FLJ12541 similar to Stra6	AF352728	1.52
complement component (3b/4b) receptor 1, including Knops blood group system	NM_000651	1.52
dystrobrevin, alpha	U46745	1.52
hypothetical protein DKFZp761O0113	NM_018409	1.52
solute carrier family 21 (prostaglandin transporter), member 2	NM_005630	1.52
striated muscle contraction regulatory protein	M96843	1.52
nucleoporin 88kD	Y08613	1.52
immunoglobulin lambda joining 3	AF234254	1.52
KIAA1685 protein	NM_018263	1.52
hypothetical protein FLJ11264	NM_018371	1.52
ets variant gene 3	L16464	1.52
Consensus includes gb:AK025026.1 /DEF=Homo sapiens cDNA: FLJ21373 fis, clone COL03214. /FEA=mRNA /DB_XREF=gi:10437461 /UG=Hs.306747 Homo sapiens cDNA: FLJ21373 fis, cl	AK025026	1.52
Consensus includes gb:X63118 /DEF=H.sapiens DNA for oestrogen receptor gene promoter /FEA=CDS /DB_XREF=gi:35159 /UG=Hs.247938 H.sapiens DNA for oestrogen receptor gene pr	X63118	1.52
paired basic amino acid cleaving enzyme (furin, membrane associated receptor protein)	NM_002569	1.52
serine/threonine kinase with Dbl- and pleckstrin homology domains	NM_007064	1.52
GLI-Kruppel family member GLI2	NM_030379	1.52
heat-shock suppressed protein 1	AF055270	1.52
CMRF35 leukocyte immunoglobulin-like receptor	NM_006678	1.52
KIAA0674 protein	AB014574	1.52
hypothetical protein DKFZp564K142 similar to implantation-associated protein	AF130104	1.52
glycoprotein M6B	N63576	1.52
Consensus includes gb:AK023909.1 /DEF=Homo sapiens cDNA FLJ13847 fis, clone THYRO1000852, highly similar to Human branched-chain amino acid aminotransferase (ECA40) mRNA. /F	AK023909	1.52
GTPase regulator associated with the focal adhesion kinase pp125	NM_015071	1.52
RAB27B, member RAS oncogene family	NM_004163	1.52
Consensus includes gb:AK027128.1 /DEF=Homo sapiens cDNA: FLJ23475 fis, clone HSI13659. /FEA=mRNA /DB_XREF=gi:10440177 /UG=Hs.288982 Homo sapiens cDNA: FLJ23475 fis, cl	AK027128	1.52
chromosome 21 open reading frame 80	NM_015227	1.52
putative acid-sensing ion channel	NM_018674	1.52
basic-helix-loop-helix-PAS protein	AL079281	1.52
cytosolic ovarian carcinoma antigen 1	NM_006375	1.52
sprouty homolog 1, antagonist of FGF signaling (Drosophila)	BF508662	1.52
myotubularin related protein 8	AL080178	1.52

gamma-glutamyl carboxylase	NM_000821	1.52
integrin, alpha 8	AI193623	1.52
chromosome 1 open reading frame 14	NM_030933	1.52
sirtuin silent mating type information regulation 2 homolog 1 (S. cerevisiae)	NM_012238	1.52
nuclear receptor coactivator 1	U59302	1.52
angiopoietin 1	U83508	1.52
Consensus includes gb:AL512697.1 /DEF=Homo sapiens mRNA; cDNA DKFZp547F134 (from clone DKFZp547F134). /FEA=mRNA /DB_XREF=gi:12224853 /UG=Hs.147587 Homo sapiens n	AL512697	1.52
5-hydroxytryptamine (serotonin) receptor 7 (adenylate cyclase-coupled)	NM_000872	1.52
golgi phosphoprotein 2	NM_016548	1.52
inositol 1,4,5-trisphosphate 3-kinase C	D38169	1.52
hypothetical protein FLJ14249 similar to HS1 binding protein 3	NM_022460	1.52
oxysterol-binding protein-related protein 1	W19983	1.52
KIAA1614 protein	AB046834	1.52
vanilloid receptor subtype 1	AF196175	1.52
hypothetical protein FLJ12673	NM_025133	1.52
KIAA0670 protein/acinus	NM_014977	1.52
adaptor-related protein complex 1, mu 2 subunit	AA910946	1.52
transcriptional adaptor 3 (ADA3, yeast homolog)-like (PCAF histone acetylase complex)	AL117487	1.52
killer cell lectin-like receptor subfamily C, member 4	U96845	1.52
flavin containing monooxygenase 4	NM_002022	1.52
retinoic acid receptor, beta	NM_000965	1.52
aryl hydrocarbon receptor	NM_001621	1.52
fibroblast growth factor receptor 2 (bacteria-expressed kinase, keratinocyte growth factor receptor, craniofacial dysostosis 1, Crouzon syndrome, Pfeiffer syndrome, Jackson-Weiss syndrome	AB030073	1.52
endoglin (Osler-Rendu-Weber syndrome 1)	BE732652	1.52
solute carrier family 7, (cationic amino acid transporter, y+ system) member 10	NM_017965	1.52
Consensus includes gb:BG231551 /FEA=EST /DB_XREF=gi:12726658 /DB_XREF=est:naf29h05.x1 /CLONE=IMAGE:4142457 /UG=Hs.74861 activated RNA polymerase II transcription cofact	BG231551	1.52
plasminogen-like	N58120	1.52
PRO1992 protein	NM_014107	1.52
Consensus includes gb:U92992.1 /DEF=Homo sapiens clone DT1P1A11 mRNA, CAG repeat region. /FEA=mRNA /DB_XREF=gi:2781410 /UG=Hs.168303 Homo sapiens clone DT1P1A11 mRI	U92992	1.52
frizzled homolog 7 (Drosophila)	AI333651	1.52
gb:AF130053.1 /DEF=Homo sapiens clone FLB4228 PRO1095 mRNA, complete cds. /FEA=mRNA /PROD=PRO1095 /DB_XREF=gi:11493412 /UG=Hs.302145 Homo sapiens clone FLB4228	AF130053	1.52
Consensus includes gb:AL050043.1 /DEF=Homo sapiens mRNA; cDNA DKFZp566M0524 (from clone DKFZp566M0524). /FEA=mRNA /DB_XREF=gi:4884285 /UG=Hs.306310 Homo sapiens	AL050043	1.52
Consensus includes gb:BE673723 /FEA=EST /DB_XREF=gi:10034264 /DB_XREF=est:7d79h10.x1 /CLONE=IMAGE:3279235 /UG=Hs.82719 Homo sapiens mRNA; cDNA DKFZp586F1822 (fr	BE673723	1.52
mitochondrial ribosomal protein L33	NM_004891	1.52
hypothetical protein FLJ20510	AF298547	1.52
aldehyde dehydrogenase 3 family, member A2	L47162	1.52
Fc fragment of IgG binding protein	NM_003890	1.52
degenerative spermatocyte homolog, lipid desaturase (Drosophila)	BC000961	1.52
sialic acid binding Ig-like lectin 7	AJ130712	1.52
hypothetical protein FLJ11292	NM_018382	1.52
actinin, alpha 1	M95178	1.52
DVS27-related protein	AB024518	1.52
diaphanous homolog 2 (Drosophila)	NM_007309	1.52
CDC14 cell division cycle 14 homolog B (S. cerevisiae)	AF064105	1.52
Ral guanine nucleotide exchange factor RalGPS1A	AF221098	1.52
killer cell immunoglobulin-like receptor, two domains, long cytoplasmic tail, 4	NM_002255	1.52
solute carrier family 4, sodium bicarbonate cotransporter, member 4	NM_003759	1.52
nuclear receptor subfamily 2, group C, member 1	M21985	1.52
dickkopf homolog 3 (Xenopus laevis)	AJ148057	1.52
calponin 1, basic, smooth muscle	NM_001299	1.52
hypothetical protein FLJ10783	NM_018217	1.52
nitric oxide synthase 1 (neuronal)	NM_000620	1.52
hypothetical protein FLJ20753	NM_017950	1.52
makorin, ring finger protein, 2	NM_014160	1.52
KIAA0985 protein	NM_014954	1.52
Consensus includes gb:BE551219 /FEA=EST /DB_XREF=gi:9792911 /DB_XREF=est:7b56b11.x1 /CLONE=IMAGE:3232221 /UG=Hs.16762 Homo sapiens mRNA; cDNA DKFZp564B2062 (fro	BE551219	1.52
Consensus includes gb:AK000995.1 /DEF=Homo sapiens cDNA FLJ10133 fis, clone HEMBA1003067. /FEA=mRNA /DB_XREF=gi:7021998 /UG=Hs.272209 Homo sapiens cDNA FLJ10133 fis	AK000995	1.52
CD1E antigen, e polypeptide	AA309511	1.52
protein phosphatase 1, regulatory (inhibitor) subunit 3C	N26005	1.52
carcinoembryonic antigen-related cell adhesion molecule 6 (non-specific cross reacting antigen)	M18728	1.52
Consensus includes gb:AU147402 /FEA=EST /DB_XREF=gi:11008923 /DB_XREF=est:AU147402 /CLONE=MAMMA1000567 /UG=Hs.287496 Homo sapiens cDNA FLJ12160 fis, clone MAMM	AU147402	1.52
KIAA0367 protein	AB002365	1.52
empty spiracles homolog 2 (Drosophila)	AI478455	1.52

melanoma antigen, family E, 1, cancer/testis specific	AL031073	1.52
small inducible cytokine A3	NM_002983	1.52
tubulin, alpha 8	NM_018943	1.52
phospholipase C, beta 4	NM_000933	1.52
KIAA1030 protein	AB028953	1.52
SHC (Src homology 2 domain-containing) transforming protein 1 pseudogene 1	Y09846	1.52
general transcription factor IIA, 1 (37kD and 19kD subunits)	NM_015859	1.52
tripartite motif-containing 3	NM_006458	1.52
sodium channel, nonvoltage-gated 1, beta (Liddle syndrome)	NM_000336	1.52
cone-rod homeobox	BE257838	1.52
activin A receptor, type IB	BC000254	1.52
Consensus includes gb:BE464799 /FEA=EST /DB_XREF=gi:9510493 /DB_XREF=est:hs87a06.s1 /CLONE=IMAGE:3144178 /UG=Hs.27342 ESTs	BE464799	1.52
cadherin 12, type 2 (N-cadherin 2)	L33477	1.52
protein (peptidyl-prolyl cis/trans isomerase) NIMA-interacting, 4 (parvulin)	BE674061	1.52
sialic acid binding Ig-like lectin 6	D86359	1.52
soggy-1 gene	NM_014419	1.52
interferon gamma receptor 1	AF056979	1.52
solute carrier family 6 (neurotransmitter transporter, glycine), member 5	AF142501	1.52
hypothetical protein MGC10902	NM_030570	1.52
fibroblast growth factor 14	NM_004115	1.52
potassium voltage-gated channel, KQT-like subfamily, member 3	NM_004519	1.52
milk fat globule-EGF factor 8 protein	BC003610	1.52
alpha thalassemia/mental retardation syndrome X-linked (RAD54 homolog, S. cerevisiae)	BC002521	1.52
mel transforming oncogene (derived from cell line NK14)- RAB8 homolog	BC002977	1.52
interleukin 1, beta	NM_000576	1.52
disabled homolog 2, mitogen-responsive phosphoprotein (Drosophila)	NM_001343	1.52
cyclin-dependent kinase-like 1 (CDC2-related kinase)	NM_004196	1.51
chromosome 9 open reading frame 7	NM_017586	1.51
KIAA1330 protein	AB037751	1.51
Consensus includes gb:AL137435.1 /DEF=Homo sapiens mRNA; cDNA DKFZp761G0924 (from clone DKFZp761G0924). /FEA=mRNA /DB_XREF=gi:6807994 /UG=Hs.274577 Homo sapiens	AL137435	1.51
dopamine receptor D2	S69899	1.51
Consensus includes gb:AF090887.1 /DEF=Homo sapiens clone HQ0085. /FEA=mRNA /DB_XREF=gi:6690154 /UG=Hs.306562 Homo sapiens clone HQ0085	AF090887	1.51
5-hydroxytryptamine (serotonin) receptor 3B	NM_006028	1.51
aldo-keto reductase family 1, member B10 (aldose reductase)	NM_020299	1.51
bromodomain and PHD finger containing, 1	NM_004634	1.51
methylmalonate-semialdehyde dehydrogenase	AF130089	1.51
Consensus includes gb:AK025177.1 /DEF=Homo sapiens cDNA: FLJ21524 fis, clone COL05921. /FEA=mRNA /DB_XREF=gi:10437640 /UG=Hs.306778 Homo sapiens cDNA: FLJ21524 fis, cl	AK025177	1.51
solute carrier family 12 (potassium/chloride transporters), member 4	AF054506	1.51
procollagen-lysine, 2-oxoglutarate 5-dioxygenase (lysine hydroxylase) 2	AJ754404	1.51
chromosome 21 open reading frame 108	NM_014825	1.51
thyroid transcription factor 1	BC006221	1.51
forkhead box I1	NM_012188	1.51
nitric oxide synthase 1 (neuronal)	U31466	1.51
magic roundabout	NM_019055	1.51
programmed death ligand 2	NM_025239	1.51
KIAA0964 protein	NM_014902	1.51
RNA binding motif protein 9	AW149379	1.51
organic cationic transporter-like 3	NM_004256	1.51
hypothetical protein	NM_016387	1.51
carbohydrate (N-acetylglucosamine 6-O) sulfotransferase 4	NM_005769	1.51
coagulation factor VII (serum prothrombin conversion accelerator)	NM_000131	1.51
protocadherin 17	NM_014459	1.51
retinal pigment epithelium-derived rhodopsin homolog	NM_006583	1.51
Consensus includes gb:AF038193.1 /DEF=Homo sapiens clone 23608 mRNA sequence. /FEA=mRNA /DB_XREF=gi:2795913 /UG=Hs.6220 Homo sapiens clone 23608 mRNA sequence	AF038193	1.51
hypothetical protein FLJ13197	NM_024614	1.51
coagulation factor II (thrombin) receptor-like 1	BE965369	1.51
chromosome 1 open reading frame 1	NM_001213	1.51
killer cell immunoglobulin-like receptor, two domains, short cytoplasmic tail, 3	NM_012313	1.51
KIAA0375 gene product	AB002373	1.51
synaptogyrin 1	NM_004711	1.51
hypothetical protein FLJ13941	NM_024848	1.51
tachykinin receptor 2	NM_001057	1.51
interphotoreceptor matrix proteoglycan 1	NM_001563	1.51

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myosin, heavy polypeptide 11, smooth muscle	NM_022844	1.51
nuclear receptor co-repressor 1	AW771910	1.51
UDP glycosyltransferase 2 family, polypeptide B15	U06641	1.51
peptidyl arginine deiminase, type V	NM_012387	1.51
synovial sarcoma, X breakpoint 3	S82471	1.51
6 antigen 4	NM_001474	1.51
Consensus includes gb:AL050328 /DEF=Human DNA sequence from clone RP11-145L22 on chromosome 6p21.32-22.2. Contains the gene for myelinoligodendrocyte glycoprotein MOG, (pa	AL050328	1.51
transforming growth factor, beta receptor III (betaglycan, 300kD)	NM_003243	1.51
CD84 antigen (leukocyte antigen)	AF054817	1.51
death effector filament-forming Ced-4-like apoptosis protein	AF310105	1.51
Consensus includes gb:AW771015 /FEA=EST /DB_XREF=gi:7703066 /DB_XREF=est:hn50d05.s1 /CLONE=IMAGE:3027081 /UG=Hs.279020 Homo sapiens mRNA; cDNA DKFZp566E0224 (h	AW771015	1.51
Consensus includes gb:BC000975.2 /DEF=Homo sapiens, clone IMAGE:3448306, mRNA, partial cds. /FEA=mRNA /PROD=Unknown (protein for IMAGE:3448306) /DB_XREF=gi:12803026 /L	BC000975	1.51
Consensus includes gb:AL031686 /DEF=Human DNA sequence from clone 981L23 on chromosome 20q12.1-13.2. Contains a Krueppel type zinc-finger protein pseudogene, a ZNF127 pseu	AL031686	1.51
Consensus includes gb:BG478428 /FEA=EST /DB_XREF=gi:13410807 /DB_XREF=est:602523839F1 /CLONE=IMAGE:4642353 /UG=Hs.326416 Homo sapiens mRNA; cDNA DKFZp564H191(	BG478428	1.51
P311 protein	U36189	1.51
hypothetical protein MGC4643	AA824369	1.51
prostaglandin E receptor 3 (subtype EP3)	D38299	1.51
interferon-induced, hepatitis C-associated microtubular aggregate protein (44kD)	NM_006417	1.51
KIAA0680 gene product	AA551142	1.51
Consensus includes gb:AA446466 /FEA=EST /DB_XREF=gi:2159131 /DB_XREF=est:zw65b12.s1 /CLONE=IMAGE:781055 /UG=Hs.193783 Human DNA sequence from clone RP13-329D4 or	AA446466	1.51
KIAA0573 protein	AI827677	1.51
pyruvate kinase, liver and RBC	M15465	1.51
Rad50-interacting protein 1	NM_021930	1.51
endothelial PAS domain protein 1	AF052094	1.51
TAF6-like RNA polymerase II, p300/CBP-associated factor (PCAF)-associated factor, 65 kD	AI005317	1.51
fibrinogen, A alpha polypeptide	NM_000508	1.51
BCL2-related ovarian killer	NM_014204	1.51
hypothetical protein	NM_012066	1.51
DnaJ (Hsp40) homolog, subfamily B, member 5	AK023253	1.51
cytochrome P450, subfamily I (dioxin-inducible), polypeptide 1 (glaucoma 3, primary infantile)	AU154504	1.51
solute carrier family 28 (sodium-coupled nucleoside transporter), member 1	U62966	1.51
actin related protein 2/3 complex, subunit 4 (20 kD)	AF019888	1.51
serine (or cysteine) proteinase inhibitor, clade E (nexin, plasminogen activator inhibitor type 1), member 1	AL574210	1.51
ectonucleoside triphosphate diphosphohydrolase 5	NM_001249	1.51
tripartite motif-containing 10	AF220122	1.51
gb:BC001957.1 /DEF=Homo sapiens, Similar to KIAA0144 gene product, clone MGC:761, mRNA, complete cds. /FEA=mRNA /PROD=Similar to KIAA0144 gene product /DB_XREF=gi:128050	BC001957	1.51
hypothetical protein	BF055311	1.51
Kruppel-like factor 2 (lung)	NM_016270	1.51
supervillin	NM_003174	1.51
RAR-related orphan receptor C	NM_005060	1.51
cholecystokinin A receptor	D85606	1.51
secreted protein of unknown function	AA679297	1.51
hypothetical protein FLJ21511	NM_025087	1.51
hypothetical protein SMAP31	AB059408	1.51
Consensus includes gb:AF003738.1 /DEF=Homo sapiens chromosome 21q22.1 anonymous mRNA sequence. /FEA=mRNA /DB_XREF=gi:2197087 /UG=Hs.268526 Homo sapiens chromoso	AF003738	1.51
heat shock 70kD protein-like 1	D85730	1.51
S-phase kinase-associated protein 2 (p45)	NM_005983	1.51
hypothetical protein FLJ12684	NM_024534	1.51
contactin 1	U07820	1.51
Fc fragment of IgE, high affinity I, receptor for; gamma polypeptide	NM_004106	1.51
fibulin 1	NM_006487	1.51
frequenin homolog (Drosophila)	NM_014286	1.51
tyrosinase-related protein 1	NM_000550	1.51
myeloid cell nuclear differentiation antigen	NM_002432	1.51
mitochondrial ribosomal protein L19	NM_014763	1.51
KIAA0820 protein	AL136712	1.51
potassium voltage-gated channel, KQT-like subfamily, member 1	AF051426	1.51
CGI-14 protein	AI829081	1.51
hypothetical protein FLJ21870	NM_023016	1.51
cadherin 5, type 2, VE-cadherin (vascular epithelium)	NM_001795	1.51
small inducible cytokine subfamily A (Cys-Cys), member 18, pulmonary and activation-regulated	AB000221	1.51
hypothetical protein FLJ20513	NM_017855	1.51
glycosylphosphatidylinositol specific phospholipase D1	NM_001503	1.51

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kinase suppressor of ras	AA971768	1.51
glycophorin B (includes Ss blood group)	U05255	1.51
alanyl (membrane) aminopeptidase (aminopeptidase N, aminopeptidase M, microsomal aminopeptidase, CD13, p150)	NM_001150	1.51
A kinase (PRKA) anchor protein 1	BC000729	1.51
potassium voltage-gated channel, Shal-related subfamily, member 3	AF120491	1.51
nuclear receptor co-repressor 2	AJ373205	1.51
Consensus includes gb:BG289001 /FEA=EST /DB_XREF=gi:13044404 /DB_XREF=est:602381262F1 /CLONE=IMAGE:4499078 /UG=Hs.101282 Homo sapiens cDNA: FLJ21238 fis, clone COL	BG289001	1.51
DEME-6 protein	BC004399	1.51
tumor necrosis factor receptor superfamily, member 11b (osteoprotegerin)	BF433902	1.51
Consensus includes gb:BE531136 /FEA=EST /DB_XREF=gi:9759781 /DB_XREF=est:601278315F1 /CLONE=IMAGE:3610539 /UG=Hs.8982 ESTs, Highly similar to KIAA1395 protein H.sapie	BE531136	1.51
Consensus includes gb:S65761.1 /DEF=anti-colorectal carcinoma heavy chain=glycoprotein CANAG-50 specific IgG1 kappa human, 19.9 hybridoma, antibody 1116NS19.9, mRNA, 1515 nt. ,	S65761	1.51
frizzled homolog 7 (Drosophila)	NM_003507	1.51
hypothetical protein similar to small G proteins, especially RAP-2A	NM_021183	1.51
NADPH oxidase 1	AF166327	1.51
serine protease inhibitor, Kazal type, 5	NM_006846	1.51
fibronectin leucine rich transmembrane protein 2	AF169676	1.51
CGI-142	AW207448	1.51
small nuclear ribonucleoprotein 70kD polypeptide (RNP antigen)	W67744	1.51
G protein-coupled receptor kinase 7	NM_017572	1.51
hypothetical protein FLJ23059	AK026712	1.51
gb:AF098641.1 /DEF=Homo sapiens CD44 isoform RC (CD44) mRNA, complete cds. /FEA=mRNA /GEN=CD44 /PROD=CD44 isoform RC /DB_XREF=gi:3832517 /UG=Hs.306278 Homo sapie	AF098641	1.51
KIAA1117 protein	AL162056	1.51
hepatocyte nuclear factor 3, beta	AB028021	1.51
tumor necrosis factor, alpha-induced protein 2	NM_006291	1.51
integrin, alpha 3 (antigen CD49C, alpha 3 subunit of VLA-3 receptor)	NM_002204	1.51
G-protein coupled receptor	AJ703188	1.51
hepatocyte nuclear factor 4, alpha	X87870	1.51
Consensus includes gb:AW072461 /FEA=EST /DB_XREF=gi:6027459 /DB_XREF=est:xa06c09.x1 /CLONE=IMAGE:2567536 /UG=Hs.169284 Homo sapiens phosphoribosylpyrophosphate sy	AW072461	1.51
Consensus includes gb:AI820796 /FEA=EST /DB_XREF=gi:5439875 /DB_XREF=est:an35h08.x5 /CLONE=IMAGE:1700703 /UG=Hs.310806 ESTs, Moderately similar to ALU7_HUMAN ALU SU	AI820796	1.51
crystallin, beta A1	NM_005208	1.51
zinc finger protein 38 (KOX 25)	AJ752257	1.50
interleukin 12 receptor, beta 2	NM_001559	1.50
phosphatidylinositol glycan, class Q	NM_004204	1.50
guanylate cyclase 1, soluble, alpha 3	AJ719730	1.50
Max-interacting protein	BE502432	1.50
a disintegrin and metalloproteinase domain 29	NM_021780	1.50
Phosphatidylglycerophosphate Synthase	NM_024419	1.50
transforming growth factor, beta 2	M19154	1.50
nuclear receptor subfamily 2, group E, member 3	NM_016346	1.50
Consensus includes gb:AL134453 /FEA=EST /DB_XREF=gi:6602640 /DB_XREF=est:DKFZp547J135_s1 /CLONE=DKFZp547J135 /UG=Hs.106890 Homo sapiens clone 23771 mRNA sequen	AL134453	1.50
hypothetical protein FLJ20234	BC000795	1.50
hypothetical protein FLJ20420	NM_017812	1.50
sialic acid binding Ig-like lectin 7	NM_016543	1.50
Consensus includes gb:AL122122.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434L098 (from clone DKFZp434L098). /FEA=mRNA /DB_XREF=gi:6102949 /UG=Hs.274559 Homo sapiens mR	AL122122	1.50
Consensus includes gb:BF434424 /FEA=EST /DB_XREF=gi:11446712 /DB_XREF=est:7o96h09.x1 /CLONE=IMAGE:3644393 /UG=Hs.107164 spectrin, beta, non-erythrocytic 1	BF434424	1.50
hypothetical protein FLJ12768	NM_025163	1.50
lymphocyte-activation gene 3	NM_002286	1.50
dermatan sulfate proteoglycan 3	NM_004950	1.50
PRO1693 protein	NM_014097	1.50
a disintegrin and metalloproteinase domain 12 (meltrin alpha)	W46291	1.50
ankyrin 3, node of Ranvier (ankyrin G)	NM_020987	1.50
proline synthetase co-transcribed (bacterial homolog)	AK021923	1.50
chondroitin sulfate proteoglycan 2 (versican)	NM_004385	1.50
Consensus includes gb:BE465829 /FEA=EST /DB_XREF=gi:9511604 /DB_XREF=est:hx80h09.x1 /CLONE=IMAGE:3194177 /UG=Hs.150539 Homo sapiens cDNA FLJ13793 fis, clone THYRO1	BE465829	1.50
protein phosphatase 1, catalytic subunit, beta isoform	NM_002709	1.50
runt-related transcription factor 1 (acute myeloid leukemia 1; aml1 oncogene)	D43967	1.50
Rag D protein	AL138717	1.50
KIAA1076 protein	AB028999	1.50
Consensus includes gb:AK025026.1 /DEF=Homo sapiens cDNA: FLJ21373 fis, clone COL03214. /FEA=mRNA /DB_XREF=gi:10437461 /UG=Hs.306747 Homo sapiens cDNA: FLJ21373 fis, cl	AK025026	1.50
amyloid beta (A4) precursor protein (protease nexin-II, Alzheimer disease)	X06989	1.50
odd-skipped-related 2A protein	AI811298	1.50
Putative prostate cancer tumor suppressor	AI884858	1.50
protein tyrosine phosphatase, receptor type, M	NM_002845	1.50

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hypothetical protein FLJ10188	NM_018017	1.50
hypothetical protein FLJ20048	NM_017640	1.50
leucine rich repeat (in FLII) interacting protein 2	NM_017724	1.50
colony stimulating factor 3 receptor (granulocyte)	NM_000760	1.50
hypothetical protein	AK025938	1.50
ankyrin repeat domain 6	BE677131	1.50
Consensus includes gb:AL523184 /FEA=EST /DB_XREF=gi:12786677 /DB_XREF=est:AL523184 /CLONE=CS0DC001YI02 (5 prime) /UG=Hs.288146 ESTs, Highly similar to KIAA0169 protein	AL523184	1.50
integrin, beta 3 (platelet glycoprotein IIIa, antigen CD61)	J02703	1.50
KIAA0934 protein	N31807	1.50
hypothetical protein FLJ12765	NM_024852	1.50
T cell receptor alpha variable 8-6	X61079	1.50
hypothetical protein	NM_013310	1.50
phospholipase A2-activating protein	AF083395	1.50
hypocretin (orexin) neuropeptide precursor	NM_001524	1.50
TAF6-like RNA polymerase II, p300/CBP-associated factor (PCAF)-associated factor, 65 kD	AI005317	1.50
olfactory receptor, family 2, subfamily W, member 1	NM_030903	1.50
NADPH oxidase, EF hand calcium-binding domain 5	NM_024505	1.50
coagulation factor XIII, A1 polypeptide	NM_000129	1.50
tumor necrosis factor receptor superfamily, member 12 (translocating chain-association membrane protein)	NM_003790	1.50
hypothetical protein MGC3047	AW888223	1.50
SRY (sex determining region Y)-box 18	NM_018419	1.50
hypothetical protein PRO1777	NM_018605	1.50
protein tyrosine phosphatase, receptor type, F	AI762627	1.50
BH-protocadherin (brain-heart)	NM_002589	1.50
fms-related tyrosine kinase 1 (vascular endothelial growth factor/vascular permeability factor receptor)	AA058828	1.50
cartilage paired-class homeoprotein 1	NM_006982	1.50
prominin-like 1 (mouse)	NM_006017	1.50
centaurin, beta 2	D26069	1.50
gap junction protein, beta 3, 31kD (connexin 31)	AF099730	1.50
diubiquitin	NM_006398	1.50
transforming growth factor, beta 3	J03241	1.50
hypothetical protein FLJ12178	NM_025134	1.50
hypothetical protein FLJ22649 similar to signal peptidase SPC22/23	NM_021928	1.50
peroxisomal trans 2-enoyl CoA reductase; putative short chain alcohol dehydrogenase	NM_018441	1.50
membrane-spanning 4-domains, subfamily A, member 3 (hematopoietic cell-specific)	L35848	1.50
hypothetical protein MGC2306	BC002557	1.50
nuclear factor of activated T-cells 5, tonicity-responsive	NM_006599	1.50
apolipoprotein B (including Ag(x) antigen)	NM_000384	1.50
jun D proto-oncogene	AI762296	1.50
salivary proline-rich protein	NM_006685	1.50
activator of CREM in testis	AL132776	1.50
met proto-oncogene (hepatocyte growth factor receptor)	AA005141	1.50
protein kinase, cGMP-dependent, type I	D45864	1.50
glycoprotein V (platelet)	L11238	1.50
fatty acid binding protein 3, muscle and heart (mammary-derived growth inhibitor)	AI041520	1.50
p8 protein (candidate of metastasis 1)	AF135266	1.50
WD repeat domain 12	NM_018256	1.50
actin-related protein 3-beta	NM_020445	1.50
gb:BC002828.1 /DEF=Homo sapiens, Similar to fetal hypothetical protein, clone MGC:3843, mRNA, complete cds. /FEA=mRNA /PROD=Similar to fetal hypothetical protein /DB_XREF=gi:12(	BC002828	1.50
Consensus includes gb:AK027151.1 /DEF=Homo sapiens cDNA: FLJ23498 fis, clone LNG02683. /FEA=mRNA /DB_XREF=gi:10440206 /UG=Hs.306904 Homo sapiens cDNA: FLJ23498 fis, cl	AK027151	1.50
hydroxyacid oxidase (glycolate oxidase) 1	NM_017545	1.50
hypothetical protein FLJ22349	NM_024821	1.50
regulatory factor X, 2 (influences HLA class II expression)	NM_000635	1.50
chromosome 11open reading frame 9	AC004770	1.50
NG3 protein	NM_030652	1.50
prodynorphin	NM_024411	1.50
Consensus includes gb:AW295295 /FEA=EST /DB_XREF=gi:6701931 /DB_XREF=est:UI-H-BI2-aht-f-12-0-UI.s1 /CLONE=IMAGE:2728079 /UG=Hs.212522 ESTs, Highly similar to B48189 so	AW295295	1.50
Consensus includes gb:AI350339 /FEA=EST /DB_XREF=gi:4087545 /DB_XREF=est:qo33g06.x1 /CLONE=IMAGE:1910362 /UG=Hs.144563 tryptophan hydroxylase (tryptophan 5-monooxyg	AI350339	1.50
DKFZP586F1018 protein	AL050090	1.50
hepatitis A virus cellular receptor 1	NM_012206	1.50
solute carrier family 9 (sodium/hydrogen exchanger), isoform 3	NM_004174	1.50
prostaglandin E synthase	NM_004878	1.50
hypothetical protein FLJ22638	NM_024839	1.50

par-3 partitioning defective 3 homolog (C. elegans)	NM_019619	1.50
tumor endothelial marker 5 precursor	AJ655903	1.50
ectonucleotide pyrophosphatase/phosphodiesterase 3	NM_005021	1.50
myelin proteolipid protein-like protein	NM_020181	1.50
pleckstrin homology domain-containing, family A (phosphoinositide binding specific) member 4	NM_020904	1.50
Consensus includes gb:AK024553.1 /DEF=Homo sapiens cDNA: FLJ20900 fis, clone ADKA03593. /FEA=mRNA /DB_XREF=gi:10436861 /UG=Hs.306687 Homo sapiens cDNA: FLJ20900 fis,	AK024553	1.50
SP110 nuclear body protein	NM_004510	1.50
class-I MHC-restricted T cell associated molecule	NM_019604	1.50
clone FLB1727	NM_016414	1.50
protein A	NM_019858	1.50
Consensus includes gb:AI207792 /FEA=EST /DB_XREF=gi:3769734 /DB_XREF=est:ao89d03.x1 /CLONE=IMAGE:1953029 /UG=Hs.306019 ESTs, Weakly similar to ALU7_HUMAN ALU SUBF	AI207792	1.50
similar to S68401 (cattle) glucose induced gene	NM_012261	1.50
hypothetical protein MGC5457	BC000988	1.50
leucine rich repeat (in FLII) interacting protein 1	BF965566	1.50
hypothetical zinc finger protein FLJ14011	NM_022103	1.50
Dystonia musculorum of mouse, human homolog of	AL049215	1.50
PRO1386 protein	NM_031269	1.50
growth hormone 1	NM_022561	1.50
ring finger protein 17	NM_031277	1.50
solute carrier family 25 (mitochondrial carrier; Graves disease autoantigen), member 16	AI827990	1.50
secretory carrier membrane protein 1	AL049223	1.50
ferrochelatase (protoporphyria)	NM_000140	1.50
haptoglobin-related protein	NM_020995	1.50
Consensus includes gb:AK024455.1 /DEF=Homo sapiens mRNA for FLJ00047 protein, partial cds. /FEA=mRNA /GEN=FLJ00047 /PROD=FLJ00047 protein /DB_XREF=gi:10440423 /UG=Hs.	AK024455	1.50
calcium channel, voltage-dependent, alpha 2/delta subunit 2	NM_006030	1.50
Consensus includes gb:AK026682.1 /DEF=Homo sapiens cDNA: FLJ23029 fis, clone LNG01883. /FEA=mRNA /DB_XREF=gi:10439589 /UG=Hs.306864 Homo sapiens cDNA: FLJ23029 fis, cl	AK026682	1.50
c-myc binding protein	NM_012333	1.50
KIAA0828 protein	AK025372	1.50
chondroitin sulfate proteoglycan BEHAB/brevican	AI205180	1.50
dymein, cytoplasmic, intermediate polypeptide 2	AF250307	1.50
similar to MYOSIN HEAVY CHAIN, CARDIAC MUSCLE ALPHA ISOFORM (MYHC-ALPHA) (M. musculus)	AK000947	1.50
phosphoinositide-3-kinase, regulatory subunit, polypeptide 2 (p85 beta)	NM_005027	1.50
Consensus includes gb:AI478300 /FEA=EST /DB_XREF=gi:4371526 /DB_XREF=est:tm39e01.x1 /CLONE=IMAGE:2160504 /UG=Hs.192789 ESTs, Weakly similar to ALU6_HUMAN ALU SUBF	AI478300	1.50
cholinergic receptor, nicotinic, alpha polypeptide 6	NM_004198	1.50
Eph82	D31661	1.50
Ser-Thr protein kinase related to the myotonic dystrophy protein kinase	NM_003607	1.50
TBP-like 1	NM_004865	1.50
KIAA0960 protein	BF447246	1.50
nephroblastoma overexpressed gene	BF440025	1.50
hypothetical protein MGC11266	NM_024322	1.50
faciogenital dysplasia (Aarskog-Scott syndrome)	NM_004463	1.50
leucine zipper, putative tumor suppressor 1	BE312985	1.50
EphA1	NM_005232	1.50
testis expressed sequence 15	NM_031271	1.50
hepatocyte nuclear factor 3, beta	AB028021	1.50
topoisomerase (DNA) III alpha	AI676092	1.50
aldehyde dehydrogenase 7 family, member A1	AU149534	1.50
hypothetical protein MGC2376	NM_023930	1.50
sialyltransferase 8 (alpha-N-acetylneuraminate: alpha-2,8-sialytransferase, GD3 synthase) A	L32867	1.50
T-box 4	NM_018488	1.50
anaplastic lymphoma kinase (Ki-1)	BF057492	1.50
cytoplasmic FMRP interacting protein 1	BC005097	1.50
melanoma antigen, family A, 3	BC000340	1.50
Rhesus blood group-associated glycoprotein	AF031549	1.50
erythropoietin	AF053356	1.50
ubiquitin specific protease 9, Y chromosome (Drosophila fat facets related)	NM_004654	1.50
phospholipid scramblase 1	NM_021105	1.49
bradykinin receptor B1	NM_000710	1.49
homolog of yeast ATP12	AF070584	1.49
NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 8 (19kD, ASHI)	AA723057	1.49
lymphocyte antigen 95 (activating NK-receptor; NK-p44)	NM_004828	1.49
peptidylarginine deiminase type I	NM_013358	1.49
A kinase (PRKA) anchor protein (yotiao) 9	NM_005751	1.49

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fibroblast growth factor 13	NM_004114	1.49
hypothetical protein MGC5395	NM_024060	1.49
related to the N terminus of tre	NM_014688	1.49
leptin receptor	U52914	1.49
KIAA1661 protein	AB051448	1.49
Consensus includes gb:AK024945.1 /DEF=Homo sapiens cDNA: FLJ21292 fis, clone COL01969. /FEA=mRNA /DB_XREF=gi:10437367 /UG=Hs.306732 Homo sapiens cDNA: FLJ21292 fis, cl	AK024945	1.49
upstream binding transcription factor, RNA polymerase I	X56687	1.49
estrogen-related receptor gamma	AF094518	1.49
KIAA1091 protein	AA349595	1.49
hypothetical protein MGC20727	AF131745	1.49
opsin 3 (encephalopsin, panopsin)	NM_014322	1.49
tumor protein 63 kDa with strong homology to p53	AF061512	1.49
ATP synthase, H+ transporting, mitochondrial F0 complex, subunit e	BC003679	1.49
Consensus includes gb:AL035587 /DEF=Human DNA sequence from clone RP3-475N16 on chromosome 6p12.3-21.2. Contains the genes for CTG4A, pre-T cell receptor alpha, a novel protei	AL035587	1.49
poly(rC) binding protein 4	BC004153	1.49
inhibin, alpha	M13981	1.49
phosphatidylserine receptor	AA351360	1.49
Consensus includes gb:AF234255.1 /DEF=Homo sapiens clone KM36 immunoglobulin light chain variable region mRNA, partial cds. /FEA=mRNA /PROD=immunoglobulin light chain variable	AF234255	1.49
hypothetical protein FLJ20555	NM_017877	1.49
dihydrolipoamide S-succinyltransferase (E2 component of 2-oxo-glutarate complex)	AC006530	1.49
Consensus includes gb:AI263044 /FEA=EST /DB_XREF=gi:3871247 /DB_XREF=est:qz29e03.x1 /CLONE=IMAGE:2028316 /UG=Hs.13438 Homo sapiens clone 24626 mRNA sequence	AI263044	1.49
KIAA0570 gene product	NM_014709	1.49
KIAA0740 gene product	AB018283	1.49
protein tyrosine phosphatase, receptor type, G	NM_002841	1.49
T-box 1	AF012130	1.49
B9 protein	NM_015681	1.49
monokine induced by gamma interferon	NM_002416	1.49
Consensus includes gb:AI346483 /FEA=EST /DB_XREF=gi:4083689 /DB_XREF=est:qp51a02.x1 /CLONE=IMAGE:1926506 /UG=Hs.132807 Homo sapiens (clone 3.8-1) MHC class I mRNA fr	AI346483	1.49
calmodulin-like 3	M58026	1.49
epidermal growth factor receptor (erythroblastic leukemia viral (v-erb-b) oncogene homolog, avian)	AF125253	1.49
hypothetical protein FLJ20079	NM_017656	1.49
lysyl oxidase-like 2	AF117949	1.49
Consensus includes gb:AK024231.1 /DEF=Homo sapiens cDNA FLJ14169 fis, clone NT2RP2002056. /FEA=mRNA /DB_XREF=gi:10436558 /UG=Hs.306670 Homo sapiens cDNA FLJ14169 fi	AK024231	1.49
protein kinase C-like 2	AI633689	1.49
Consensus includes gb:AV761014 /FEA=EST /DB_XREF=gi:10918862 /DB_XREF=est:AV761014 /CLONE=MDSCCF09 /UG=Hs.291972 ESTs, Moderately similar to SC14_HUMAN SEC14-LIKE	AV761014	1.49
survival motor neuron pseudogene	U91640	1.49
hypothetical protein FLJ20689	NM_017972	1.49
complement component 1, s subcomponent	M18767	1.49
kinesin family member C3	BC001211	1.49
B-cell CLL/lymphoma 2	NM_000657	1.49
adaptor-related protein complex 4, epsilon 1 subunit	AB030653	1.49
RAR-related orphan receptor B	NM_006914	1.49
pregnancy specific beta-1-glycoprotein 2	NM_031246	1.49
calcitonin gene-related peptide-receptor component protein	AU154853	1.49
zinc finger protein 237	NM_014242	1.49
malic enzyme 1, NADP(+)-dependent, cytosolic	L34035	1.49
serine (or cysteine) proteinase inhibitor, clade B (ovalbumin), member 3	AB046400	1.49
par-3 partitioning defective 3 homolog (C. elegans)	AW452651	1.49
cell division cycle 27	AA166684	1.49
Consensus includes gb:AK000834.1 /DEF=Homo sapiens cDNA FLJ20827 fis, clone ADKA03543. /FEA=mRNA /DB_XREF=gi:7021152 /UG=Hs.306409 Homo sapiens cDNA FLJ20827 fis, clo	AK000834	1.49
paternally expressed 3	AL042588	1.49
sema domain, transmembrane domain (TM), and cytoplasmic domain, (semaphorin) 6A	NM_020796	1.49
gb:AB058895.1 /DEF=Homo sapiens GHRHRpsv mRNA for growth hormone releasing hormone receptor pituitary splice variant, complete cds. /FEA=mRNA /GEN=GHRHRpsv /PROD=growth h	AB058895	1.49
double homeobox, 3	NM_012148	1.49
cystatin C (amyloid angiopathy and cerebral hemorrhage)	NM_000099	1.49
phosphoinositide-specific phospholipase C-beta 1	AY004175	1.49
v-rel reticuloendotheliosis viral oncogene homolog (avian)	NM_002908	1.49
likely homolog of rat kinase D-interacting substance of 220 kDa	AB033076	1.49
gb:NM_016064.1 /DEF=Homo sapiens CGI-131 protein (LOC51648), mRNA. /FEA=mRNA /GEN=LOC51648 /PROD=CGI-131 protein /DB_XREF=gi:7706344 /UG=Hs.109733 CGI-131 protei	NM_016064	1.49
KIAA0605 gene product	NM_014694	1.49
aldehyde dehydrogenase 3 family, member A1	NM_000691	1.49
hypothetical protein PRO1787	NM_018606	1.49
Consensus includes gb:AI186064 /FEA=EST /DB_XREF=gi:3736702 /DB_XREF=est:qe50h05.x1 /CLONE=IMAGE:1742457 /UG=Hs.170212 ESTs, Highly similar to b-TRCP variant E3RS-Ikapp	AI186064	1.49

ecotropic viral integration site 5	NM_005665	1.49
lactoperoxidase	U39573	1.49
similar to DNA-directed RNA polymerase I (135 kDa)	NM_019014	1.49
solute carrier family 9 (sodium/hydrogen exchanger), isoform 2	NM_003048	1.49
Thy-1 cell surface antigen	AA218868	1.49
sialophorin (gpL115, leukosialin, CD43)	X60502	1.49
popeye protein 3	NM_022361	1.49
Cluster Incl. AI738452:wi32a07.x1 Homo sapiens cDNA, 3' end /clone=IMAGE-2391924 /clone_end=3' /gb=AI738452 /gi=5100433 /ug=Hs.203286 /len=533	AI738452	1.49
protease inhibitor 15	NM_015886	1.49
CDV-1 protein	NM_014055	1.49
Consensus includes gb:AA631242 /FEA=EST /DB_XREF=gi:2553853 /DB_XREF=est:nq81b04.s1 /CLONE=IMAGE:1158703 /UG=Hs.21349 ESTs, Moderately similar to RB15 RAT RAS-RELATI	AA631242	1.49
hypothetical protein FLJ10381	NM_018080	1.49
hypothetical protein FLJ20127	NM_017678	1.49
Consensus includes gb:W26966 /FEA=EST /DB_XREF=gi:1306194 /DB_XREF=est:16h5 /UG=Hs.293838 ESTs	W26966	1.49
keratin, hair, acidic, 6	NM_003771	1.49
opsin 1 (cone pigments), medium-wave-sensitive (color blindness, deutan)	NM_000513	1.49
guanine nucleotide binding protein (G protein), alpha 11 (Gq class)	M69013	1.49
ATPase, H+ transporting, lysosomal (vacuolar proton pump) non-catalytic accessory protein 1B	NM_020632	1.49
endothelial differentiation, sphingolipid G-protein-coupled receptor, 1	NM_001400	1.49
hyaluronoglucosaminidase 2	NM_003773	1.49
Dvl-binding protein IDAX (inhibition of the Dvl and Axin complex)	NM_025212	1.49
differentiation-related protein dif13	NM_016380	1.49
Ets2 repressor factor	NM_006494	1.49
cyclic nucleotide gated channel beta 1	U58837	1.49
ryanodine receptor 3	NM_001036	1.49
prostatic binding protein	NM_002567	1.49
proprotein convertase subtilisin/kexin type 2	NM_002594	1.49
AT2 receptor-interacting protein 1	AL096842	1.49
HT021	NM_020685	1.49
kinesin family member 3C	AF035621	1.49
AKAP-binding sperm protein ropporin	NM_017578	1.49
tenascin XB	X71923	1.49
ATPase, Class VI, type 11A	AL161996	1.49
DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 18 (Myc-regulated)	X98743	1.49
neurotrophic tyrosine kinase, receptor, type 3	AI613045	1.49
arylsulfatase B	NM_000046	1.49
Consensus includes gb:AK023365.1 /DEF=Homo sapiens cDNA FLJ13303 fis, clone OVARC1001372, highly similar to Homo sapiens liprin-alpha4 mRNA. /FEA=mRNA /DB_XREF=gi:104352	AK023365	1.49
nuclear receptor subfamily 5, group A, member 1	U76388	1.49
xylulokinase (H. influenzae) homolog	NM_005108	1.49
dopachrome tautomerase (dopachrome delta-isomerase, tyrosine-related protein 2)	NM_001922	1.49
tissue inhibitor of metalloproteinase 1 (erythroid potentiating activity, collagenase inhibitor)	NM_003254	1.49
regulator of mitotic spindle assembly 1	NM_002932	1.49
PR domain containing 14	NM_024504	1.49
Consensus includes gb:BF509345 /FEA=EST /DB_XREF=gi:11592709 /DB_XREF=est:UI-H-BI4-aoz-c-11-0-UI.s1 /CLONE=IMAGE:3086229 /UG=Hs.6996 ESTs	BF509345	1.49
neurotrophic tyrosine kinase, receptor, type 2	BF674712	1.49
adhesion glycoprotein	NM_006566	1.49
SET binding protein 1	NM_015559	1.49
gb:U48722.1 /DEF=Human epidermal growth factor receptor precursor (EGFR) mRNA, complete cds. /FEA=mRNA /PROD=epidermal growth factor receptor precursor /DB_XREF=gi:162854	U48722	1.49
vitamin D (1,25- dihydroxyvitamin D3) receptor	AA454701	1.49
gb:AF116689.1 /DEF=Homo sapiens PRO2168 mRNA, complete cds. /FEA=mRNA /PROD=PRO2168 /DB_XREF=gi:7959876 /UG=Hs.249715 Homo sapiens PRO2168 mRNA, complete cds /	AF116689	1.49
four jointed box 1 (Drosophila)	NM_014344	1.49
neuropilin 1	AF280547	1.49
ELK4, ETS-domain protein (SRF accessory protein 1)	NM_001973	1.49
hypothetical gene C6018	AL049786	1.49
retinoic acid receptor-beta associated open reading frame	NM_015854	1.49
homeo box D1	NM_024501	1.49
Consensus includes gb:AK026465.1 /DEF=Homo sapiens cDNA: FLJ22812 fis, clone KAIA2955. /FEA=mRNA /DB_XREF=gi:10439335 /UG=Hs.306853 Homo sapiens cDNA: FLJ22812 fis, cle	AK026465	1.49
growth factor receptor-bound protein 10	D86962	1.49
claudin 10	NM_006984	1.49
sodium channel, voltage-gated, type II, beta polypeptide	U87555	1.49
ret finger protein-like 2	NM_006605	1.49
hypothetical protein FLJ10998	NM_018294	1.49
GRO3 oncogene	NM_002090	1.49

Consensus includes gb:AW301235 /FEA=EST /DB_XREF=gi:6710912 /DB_XREF=est:xs69c06.x1 /CLONE=IMAGE:2774890 /UG=Hs.66185 Homo Sapiens mRNA, partial cDNA sequence from	AW301235	1.49
Consensus includes gb:AW851559 /FEA=EST /DB_XREF=gi:7947076 /DB_XREF=est:IL3-CT0221-040400-107-E09 /UG=Hs.202695 Human soluble CD44 (CD44) mRNA, with exon v9 extens	AW851559	1.48
cathepsin F	NM_003793	1.48
calcitonin/calcitonin-related polypeptide, alpha	M26095	1.48
thrombomodulin	NM_000361	1.48
T cell receptor gamma locus	M27331	1.48
NPC1 (Niemann-Pick disease, type C1, gene)-like 1	NM_013389	1.48
chondroitin sulfate proteoglycan 2 (versican)	D32039	1.48
neurogenin 1	NM_006161	1.48
pleckstrin	AI433595	1.48
transforming, acidic coiled-coil containing protein 1	AB029026	1.48
Consensus includes gb:AK000786.1 /DEF=Homo sapiens cDNA FLJ20779 fis, clone COL05077. /FEA=mRNA /DB_XREF=gi:7021088 /UG=Hs.321583 Homo sapiens cDNA FLJ20779 fis, clon	AK000786	1.48
glioma-associated oncogene homolog (zinc finger protein)	NM_005269	1.48
matrix metalloproteinase 17 (membrane-inserted)	NM_016155	1.48
lipase, hepatic	NM_000236	1.48
polycystic kidney disease 1 (autosomal dominant)	L39891	1.48
olfactory receptor, family 52, subfamily A, member 1	NM_012375	1.48
microtubule-associated protein tau	J03778	1.48
Consensus includes gb:AW836210 /FEA=EST /DB_XREF=gi:7930184 /DB_XREF=est:RC3-LT0023-200100-012-e08 /UG=Hs.278648 Homo sapiens cDNA FLJ14085 fis, clone HEMBB10025:	AW836210	1.48
Consensus includes gb:AL353942.1 /DEF=Homo sapiens mRNA; cDNA DKFZp761L23121 (from clone DKFZp761L23121). /FEA=mRNA /DB_XREF=gi:7669981 /UG=Hs.306504 Homo sapier	AL353942	1.48
putative G protein coupled receptor	NM_007223	1.48
pro-melanin-concentrating hormone-like 1	S64288	1.48
GTP-binding protein homologous to Saccharomyces cerevisiae SEC4	NM_006822	1.48
interferon, alpha 2	M54886	1.48
caspase 2, apoptosis-related cysteine protease (neural precursor cell expressed, developmentally down-regulated 2)	BC002427	1.48
lysophosphatidic acid acyltransferase-delta	NM_020133	1.48
Oyster Incl, AI806793:wf15d05.x1 Homo sapiens cDNA, 3' end /clone=IMAGE-2350665 /clone_end=3' /gb=AI806793 /gi=5393359 /ug=Hs.239450 /len=555	AI806793	1.48
phosphodiesterase 3A, cGMP-inhibited	U36798	1.48
mitogen-activated protein kinase kinase kinase 2	AF239798	1.48
Consensus includes gb:AL359553 /DEF=Human DNA sequence from clone RP5-871G17 on chromosome 1. Contains ESTs, STSs and GSSs. Contains the HAO2 gene for hydroxyacid oxidase 2	AL359553	1.48
CD44 antigen (homing function and Indian blood group system)	BC004372	1.48
glucagon-like peptide 1 receptor	L23503	1.48
tropomodulin 2 (neuronal)	NM_014548	1.48
interleukin 2	NM_000586	1.48
Kruppel-associated box protein	NM_015877	1.48
reserved	NM_020646	1.48
Consensus includes gb:Z21818 /DEF=H.sapiens carcinoembryonic antigen gene /FEA=mRNA /DB_XREF=gi:437783 /UG=Hs.166040 H.sapiens carcinoembryonic antigen gene	Z21818	1.48
CD47 antigen (Rh-related antigen, integrin-associated signal transducer)	BG230614	1.48
forkhead box J1	NM_001454	1.48
pleiotrophin (heparin binding growth factor 8, neurite growth-promoting factor 1)	BC005916	1.48
a disintegrin and metalloproteinase domain 28	NM_021777	1.48
bromodomain-containing 1	NM_014577	1.48
hydroxysteroid (11-beta) dehydrogenase 2	NM_000196	1.48
T-box 2	U28049	1.48
Consensus includes gb:AL137715.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434K0423 (from clone DKFZp434K0423); partial cds. /FEA=mRNA /GEN=DKFZp434K0423 /PROD=hypothetica	AL137715	1.48
hypothetical protein FLJ21120	NM_024767	1.48
hypothetical protein EST00098	AK022885	1.48
calcium channel, voltage-dependent, gamma subunit 3	NM_006539	1.48
reticulocalbin 2, EF-hand calcium binding domain	NM_002902	1.48
coagulation factor II (thrombin)	NM_000506	1.48
acidic epididymal glycoprotein-like 1	NM_001131	1.48
four and a half LIM domains 3	NM_004468	1.48
KIAA0328 protein	AI825212	1.48
Kelch motif containing protein	BC005253	1.48
membrane protein, palmitoylated 1 (55kD)	NM_002436	1.48
annexin A9	BC005830	1.48
Consensus includes gb:BC004857.1 /DEF=Homo sapiens, clone IMAGE:3690478, mRNA, partial cds. /FEA=mRNA /PROD=Unknown (protein for IMAGE:3690478) /DB_XREF=gi:13436058 /L	BC004857	1.48
cathepsin K (pycnodysostosis)	NM_000396	1.48
intersectin 1 (SH3 domain protein)	AF114488	1.48
reticulocalbin 1, EF-hand calcium binding domain	NM_002901	1.48
PR domain containing 9	NM_020227	1.48
lethal (3) malignant brain tumor l(3)mbt protein (Drosophila) homolog	U89358	1.48
tumor necrosis factor receptor superfamily, member 12 (translocating chain-association membrane protein)	U72763	1.48

hypothetical protein FLJ10312	NM_030672	1.48
KIAA0768 protein	R50822	1.48
Consensus includes gb:BE786598 /FEA=EST /DB_XREF=gi:10207796 /DB_XREF=est:601478824F1 /CLONE=IMAGE:3881717 /UG=Hs.101474 KIAA0807 protein	BE786598	1.48
neuronal Shc adaptor homolog	AV705938	1.48
hypothetical protein MGC2477	BC001434	1.48
ribonuclease P (38kD)	AL134489	1.48
myosin VI	AA877789	1.48
tenascin XB	AF086641	1.48
giant axonal neuropathy (gigaxonin)	NM_022041	1.48
proline-serine-threonine phosphatase interacting protein 1	AF038602	1.48
hypothetical protein FLJ22202	NM_024883	1.48
hypothetical protein FLJ13639	NM_024705	1.48
integrin, beta 3 (platelet glycoprotein IIIa, antigen CD61)	NM_000212	1.48
hypothetical protein FLJ22087	AA555088	1.48
zinc finger protein 331; zinc finger protein 463	NM_018555	1.48
interleukin 3 receptor, alpha (low affinity)	NM_002183	1.48
hypothetical protein FLJ20276	NM_017738	1.48
tripartite motif-containing 31	NM_007028	1.48
Consensus includes gb:Z21967.1 /DEF=H.sapiens partial cDNA for homologue of mPOU homeobox protein. /FEA=mRNA /PROD=mPOU homeobox protein /DB_XREF=gi:437875 /UG=Hs.28	Z21967	1.48
hypothetical protein FLJ23392	NM_024784	1.48
pregnancy specific beta-1-glycoprotein 3	NM_021016	1.48
ATPase, Na+/K+ transporting, alpha 2 (+) polypeptide	AW440492	1.48
hypothetical protein FLJ20309	NM_017759	1.48
ABO blood group (transferase A, alpha 1-3-N-acetylgalactosaminyltransferase; transferase B, alpha 1-3-galactosyltransferase)	U15197	1.48
thymosin, beta 4, Y chromosome	NM_004202	1.48
KIAA0793 gene product	NM_014808	1.48
src family associated phosphoprotein 1	NM_003726	1.48
linked to Surfeit genes in Fugu rubripes 2; LSFR2 gene 2	AI684664	1.48
suppression of tumorigenicity 7	NM_021908	1.48
zinc finger protein 6 (CMPX1)	NM_021998	1.48
pre-B-cell leukemia transcription factor 2	BC003111	1.48
hypothetical protein FLJ10595	NM_020117	1.48
phosphodiesterase 6A, cGMP-specific, rod, alpha	NM_000440	1.48
aldolase B, fructose-bisphosphate	AV652403	1.48
hypothetical protein FLJ13340	NM_025085	1.48
solute carrier family 8 (sodium/calcium exchanger), member 1	AF128524	1.48
hypothetical protein DKFZp434E169	AL136902	1.48
glycerol kinase pseudogene 1	X78713	1.48
collagen, type IV, alpha 5 (Alport syndrome)	AW052179	1.48
retinoblastoma binding protein 1	NM_002892	1.48
G-protein coupled receptor	NM_007369	1.48
TBC1 (tre-2/USP6, BUB2, cdc16) domain family, member 1	BE675153	1.48
synovial sarcoma, X breakpoint 3	BF184274	1.48
cholinergic receptor, muscarinic 5	NM_012125	1.48
decidual protein induced by progesterone	AL136653	1.48
hypothetical protein PRO0813	NM_018593	1.48
Consensus includes gb:AF052146.1 /DEF=Homo sapiens clone 24653 mRNA sequence. /FEA=mRNA /DB_XREF=gi:3360455 /UG=Hs.155085 Homo sapiens clone 24653 mRNA sequence	AF052146	1.48
cytochrome P450, subfamily IIE (ethanol-inducible)	AV650427	1.48
transmembrane 4 superfamily member 1	M90657	1.48
hypothetical protein FLJ10701	NM_018183	1.48
scaffold attachment factor B	AI761858	1.48
clusterin-like 1 (retinal)	NM_014410	1.48
erythroid differentiation-related factor	NM_016633	1.48
beaded filament structural protein 1, filensin	NM_001195	1.48
Consensus includes gb:AL583520 /FEA=EST /DB_XREF=gi:12952562 /DB_XREF=est:AL583520 /CLONE=CS0DC024YE13 (5 prime) /UG=Hs.182183 Homo sapiens mRNA for caldesmon, 3	AL583520	1.48
Consensus includes gb:Z84717.1 /DEF=Human DNA sequence from cDNA 16pHQG;19 from chromosome 16p13.3. /FEA=mRNA /DB_XREF=gi:1817571 /UG=Hs.306284 Human DNA sequen	Z84717	1.48
KIAA0534 protein	AI817331	1.48
scieilin	NM_003843	1.48
Consensus includes gb:X63749 /DEF=H.sapiens DNA for Rod transducin alpha-subunit /FEA=mRNA /DB_XREF=gi:37049 /UG=Hs.51147 guanine nucleotide binding protein (G protein), alpl	X63749	1.48
KIAA0756 protein	AI821777	1.48
ankyrin repeat and SOCS box-containing 4	AC003079	1.48
zinc finger protein, X-linked	R51161	1.48
CD5 antigen (p56-62)	NM_014207	1.48

Consensus includes gb:AL117549.1 /DEF=Homo sapiens mRNA; cDNA DKFZp586H0123 (from clone DKFZp586H0123). /FEA=mRNA /DB_XREF=gi:5912083 /UG=Hs.268496 Homo sapiens	AL117549	1.48
protease, serine, 12 (neurotrypsin, motopsin)	AI810767	1.48
alpha2,8-sialyltransferase	NM_013305	1.48
hypothetical protein MGC2601	NM_024042	1.48
achaete-scute complex-like 1 (Drosophila)	AW950513	1.48
SRY (sex determining region Y)-box 13	AF116571	1.48
Consensus includes gb:BG025063 /FEA=EST /DB_XREF=gi:12411278 /DB_XREF=est:602276406F1 /CLONE=IMAGE:4364304 /UG=Hs.44888 ESTs	BG025063	1.48
serine/threonine kinase 38	AF034187	1.48
SRY (sex determining region Y)-box 13	AF083105	1.48
paired basic amino acid cleaving system 4	D28514	1.48
Consensus includes gb:AF043583.1 /DEF=Homo sapiens clone ASMneg1-b3 immunoglobulin lambda chain VI region, (IGL) mRNA, partial cds. /FEA=mRNA /GEN=IGL /PROD=immunoglobul	AF043583	1.48
hypothetical protein DKFZp761B2423	AI538172	1.48
carcinoembryonic antigen-related cell adhesion molecule 1 (biliary glycoprotein)	X16354	1.48
Consensus includes gb:AW968210 /FEA=EST /DB_XREF=gi:8158170 /DB_XREF=est:EST380405 /UG=Hs.293957 ESTs, Weakly similar to ALUC_HUMAN !!!! ALU CLASS C WARNING ENTRY !!!	AW968210	1.48
fibronectin leucine rich transmembrane protein 1	AF169675	1.48
CD47 antigen (Rh-related antigen, integrin-associated signal transducer)	BF693956	1.48
solute carrier family 2 (facilitated glucose transporter), member 3	AL110298	1.48
interleukin 1 receptor antagonist	BE563442	1.48
RAD23 homolog B (S. cerevisiae)	T93562	1.48
allograft inflammatory factor 1	AF299327	1.48
Wolf-Hirschhorn syndrome candidate 1-like 1	NM_017778	1.47
engrailed homolog 1	NM_001426	1.47
serine/threonine kinase 3 (STE20 homolog, yeast)	NM_006281	1.47
FAT tumor suppressor homolog 2 (Drosophila)	NM_001447	1.47
DKFZP564L0862 protein	NM_024087	1.47
PRO0233 protein	NM_014121	1.47
gb:NM_030977.1 /DEF=Homo sapiens keratin associated protein KRTAP2.1A (KRTAP2.1A), mRNA. /FEA=mRNA /GEN=KRTAP2.1A /PROD=keratin associated protein KRTAP2.1A /DB_XREF=	NM_030977	1.47
killer cell lectin-like receptor subfamily F, member 1	NM_016523	1.47
solute carrier family 13 (sodium-dependent dicarboxylate transporter), member 3	NM_022829	1.47
chemokine (C-C motif) receptor 7	NM_001838	1.47
high-mobility group (nonhistone chromosomal) protein 4	NM_005342	1.47
PH domain containing protein in retina 1	AF081583	1.47
gb:AL136548.1 /DEF=Homo sapiens mRNA; cDNA DKFZp761G18121 (from clone DKFZp761G18121); complete cds. /FEA=mRNA /GEN=DKFZp761G18121 /PROD=hypothetical protein /DB_	AL136548	1.47
syntaxin 5A	NM_003164	1.47
MBP protein	NM_016586	1.47
hypothetical protein FLJ12552	AK024318	1.47
Nijmegen breakage syndrome 1 (nibrin)	NM_002485	1.47
hypothetical protein	AV700415	1.47
trans-prenyltransferase	NM_014317	1.47
transducin (beta)-like 2	AK026529	1.47
retinoblastoma 1 (including osteosarcoma)	M19701	1.47
hypothetical protein FLJ23233	NM_024691	1.47
glioma pathogenesis-related protein	NM_006851	1.47
Consensus includes gb:AU147799 /FEA=EST /DB_XREF=gi:11009320 /DB_XREF=est:AU147799 /CLONE=MAMMA1001744 /UG=Hs.288467 Homo sapiens cDNA FLJ12280 fis, clone MAMM	AU147799	1.47
taste receptor, type 2, member 10	NM_023921	1.47
nudix (nucleoside diphosphate linked moiety X)-type motif 4	AF191654	1.47
sine oculis homeobox homolog 1 (Drosophila)	NM_005982	1.47
Consensus includes gb:AK024445.1 /DEF=Homo sapiens mRNA for FLJ00035 protein, partial cds. /FEA=mRNA /GEN=FLJ00035 /PROD=FLJ00035 protein /DB_XREF=gi:10440403 /UG=Hs.	AK024445	1.47
hypothetical protein FLJ13576	NM_022484	1.47
four and a half LIM domains 2	NM_001450	1.47
met proto-oncogene (hepatocyte growth factor receptor)	BG170541	1.47
melanoma antigen, family E, 1, cancer/testis specific	NM_016249	1.47
Consensus includes gb:Z34282.1 /DEF=H.sapiens (MAR11) MUC5AC mRNA for mucin (partial). /FEA=mRNA /GEN=MUC5AC /PROD=mucin /DB_XREF=gi:563384 /UG=Hs.247953 H.sapiens	Z34282	1.47
ADP-ribosylation factor-like 7	AW450363	1.47
hypothetical protein FLJ12983	NM_024856	1.47
corticotropin releasing hormone	BC002599	1.47
Consensus includes gb:AA570453 /FEA=EST /DB_XREF=gi:2344433 /DB_XREF=est:nk62f11.s1 /CLONE=IMAGE:1018125 /UG=Hs.10323 Homo sapiens mRNA from chromosome 5q31-33 r	AA570453	1.47
collagen, type V, alpha 1	M76729	1.47
hypothetical protein FLJ13236	NM_024902	1.47
DKFZP434F1735 protein	NM_015590	1.47
DnaJ (Hsp40) homolog, subfamily B, member 5	NM_012266	1.47
chloride intracellular channel 2	AI768628	1.47
plasmolipin	AL020989	1.47

DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 28	AI922937	1.41
ras homolog gene family, member E	BG054844	1.41
potassium inwardly-rectifying channel, subfamily J, member 16	NM_018658	1.41
KIAA0172 protein	D79994	1.41
Bacillus subtilis /REF=M24537 /DEF=B subtilis pheB, pheA genes corresponding to nucleotides 2897-3200 of M24537 /LEN=1409 (-5, -M, -3 represent transcript regions 5 prime, Middle,	M24537	1.41
glutamate receptor, ionotropic, N-methyl D-aspartate 1	NM_007327	1.41
hypothetical protein FLJ11275	NM_018376	1.41
progesterone-induced blocking factor 1	NM_006346	1.41
LOC88523	U50529	1.41
zinc finger protein 133 (clone pHZ-13)	U09366	1.41
colony stimulating factor 1 receptor, formerly McDonough feline sarcoma viral (v-fms) oncogene homolog	NM_005211	1.41
ADP-ribosyltransferase 3	U47054	1.41
Consensus includes gb:L34409.1 /DEF=Homo Sapiens (clone B3B3E13) chromosome 4p16.3 DNA fragment. /FEA=mRNA /DB_XREF=gi:600517 /UG=Hs.94799 Homo Sapiens (clone B3B3I	L34409	1.41
DKFZP566K023 protein	NM_015485	1.41
dual specificity phosphatase 3 (vaccinia virus phosphatase VH1-related)	BC002682	1.41
Consensus includes gb:AL390145.1 /DEF=Homo sapiens mRNA; cDNA DKFZp762C115 (from clone DKFZp762C115). /FEA=mRNA /DB_XREF=gi:9368878 /UG=Hs.284330 Homo sapiens mR	AL390145	1.41
Dnaj (Hsp40) homolog, subfamily C, member 3	NM_006260	1.41
Consensus includes gb:AK025352.1 /DEF=Homo sapiens cDNA: FLJ21699 fis, clone COL09829. /FEA=mRNA /DB_XREF=gi:10437851 /UG=Hs.306804 Homo sapiens cDNA: FLJ21699 fis, cl	AK025352	1.41
fibronectin leucine rich transmembrane protein 2	NM_013231	1.41
solute carrier family 16 (monocarboxylic acid transporters), member 5	AA705628	1.41
hypothetical protein FLJ14260	NM_025027	1.41
Rap1 guanine-nucleotide-exchange factor directly activated by cAMP	U78168	1.41
kidney and liver proline oxidase 1	U95090	1.41
Consensus includes gb:AI004009 /FEA=EST /DB_XREF=gi:3213519 /DB_XREF=est:ou03d10.x1 /CLONE=IMAGE:1625203 /UG=Hs.130816 ESTs, Moderately similar to ALU7_HUMAN ALU SI	AI004009	1.41
transient receptor potential channel 3	NM_003305	1.41
phosphodiesterase 4D interacting protein (myomegalin)	AI950145	1.41
keratin, hair, acidic, 5	X90763	1.41
atrophin-1 interacting protein 1; activin receptor interacting protein 1	AB014605	1.41
interleukin 16 (lymphocyte chemoattractant factor)	M90391	1.41
dynein, axonemal, intermediate polypeptide, 1	NM_012144	1.41
uronyl 2-sulfotransferase	NM_005715	1.41
Consensus includes gb:AW971248 /FEA=EST /DB_XREF=gi:8161093 /DB_XREF=est:EST383337 /UG=Hs.291289 ESTs, Weakly similar to ALU1_HUMAN ALU SUBFAMILY J SEQUENCE CONTAI	AW971248	1.41
ubiquitin associated protein	AL136733	1.41
septin 6	AL568374	1.41
EH-domain containing 3	NM_014600	1.41
hypothetical protein FLJ22184	NM_025094	1.41
hypothetical protein FLJ20034	NM_017630	1.41
GDP-fucose transporter 1	NM_018389	1.41
protease, serine, 12 (neurotrypsin, motopsin)	NM_003619	1.41
Chediak-Higashi syndrome 1	U70064	1.41
fetuin B	AB017551	1.41
integrin, beta 5	NM_002213	1.41
gb:BC004241.1 /DEF=Homo sapiens, Similar to laminin, alpha 4, clone MGC:10389, mRNA, complete cds. /FEA=mRNA /PROD=Similar to laminin, alpha 4 /DB_XREF=gi:13278992 /UG=Hs.	BC004241	1.41
Consensus includes gb:U79265.1 /DEF=Human clone 23614 mRNA sequence. /FEA=mRNA /DB_XREF=gi:1710224 /UG=Hs.91375 Human clone 23614 mRNA sequence	U79265	1.41
MCM3 minichromosome maintenance deficient 3 (S. cerevisiae) associated protein	NM_003906	1.41
bromodomain adjacent to zinc finger domain, 2A	AA788652	1.41
hypothetical protein similar to small G proteins, especially RAP-2A	NM_021183	1.41
helicase-moi	NM_030621	1.41
regulator of G-protein signalling 5	AI183997	1.41
NEL-like 2 (chicken)	NM_006159	1.41
Consensus includes gb:X97875 /DEF=H.sapiens EP4 prostaglandin receptor pseudogene c16g7 /FEA=CDS /DB_XREF=gi:1359737 /UG=Hs.247972 H.sapiens EP4 prostaglandin receptor p:	X97875	1.41
cadherin 8, type 2	T65123	1.41
hypothetical protein	NM_013341	1.41
CD84 antigen (leukocyte antigen)	AF054818	1.41
lysyl oxidase	NM_002317	1.41
protein tyrosine phosphatase, receptor type, B	NM_002837	1.41
GLI-Kruppel family member GLI3 (Greig cephalopolysyndactyly syndrome)	NM_000168	1.41
cytochrome P450, subfamily IIC (mephenytoin 4-hydroxylase), polypeptide 9	M15331	1.41
syndecan 2 (heparan sulfate proteoglycan 1, cell surface-associated, fibroglycan)	BE048514	1.41
Cockayne syndrome 1 (classical)	NM_000082	1.41
apolipoprotein B mRNA editing enzyme, catalytic polypeptide-like 2	NM_006789	1.41
phosphorylase, glycogen; muscle (McArdle syndrome, glycogen storage disease type V)	NM_005609	1.41
Consensus includes gb:AK024915.1 /DEF=Homo sapiens cDNA: FLJ21262 fis, clone COL01494. /FEA=mRNA /DB_XREF=gi:10437330 /UG=Hs.306727 Homo sapiens cDNA: FLJ21262 fis, cl	AK024915	1.41

serine/threonine protein kinase MASK	NM_016542	1.41
KIAA1048 protein	NM_014911	1.41
chimerin (chimaerin) 2	AK026415	1.41
IL2-inducible T-cell kinase	D13720	1.41
hypothetical protein FLJ21916	NM_023112	1.41
collagen, type I, alpha 2	NM_000089	1.41
interferon induced transmembrane protein 3 (1-8U)	BF338947	1.41
6-phosphofructo-2-kinase/fructose-2,6-biphosphatase 1	NM_002625	1.41
DKFZP586N0721 protein	BF971416	1.41
interleukin 7 receptor	NM_002185	1.41
nuclear transport factor 2 (placental protein 15)	NM_005796	1.41
AXL receptor tyrosine kinase	AI467916	1.41
dachshund homolog (Drosophila)	NM_004392	1.41
solute carrier family 8 (sodium-calcium exchanger), member 2	AI127885	1.41
putative dimethyladenosine transferase	AF091078	1.41
ATPase, Class V, type 10B	AW006935	1.41
Consensus includes gb:AI251844.1 /DEF=Homo sapiens partial mRNA for MOZCBP chimeric transcript type II. /FEA=mRNA /GEN=MOZCBP /PROD=MOZCBP /DB_XREF=gi:9368433 /UG=Hs.	AI251844	1.41
hypothetical protein FLJ20515	NM_017857	1.41
hypothetical protein	AA160474	1.41
olfactory receptor, family 2, subfamily H, member 3	NM_007160	1.41
E3 ubiquitin ligase SMURF1	AB046845	1.41
mab-21-like 2 (C. elegans)	AF262032	1.41
Consensus includes gb:AU144791 /FEA=EST /DB_XREF=gi:11006312 /DB_XREF=est:AU144791 /CLONE=HEMBA1002971 /UG=Hs.274237 Homo sapiens cDNA FLJ10126 fis, clone HEMBA	AU144791	1.41
solute carrier family 4, sodium bicarbonate cotransporter, member 4	AF069510	1.41
neural precursor cell expressed, developmentally down-regulated 4-like	AI357376	1.41
glutamate decarboxylase 2 (pancreatic islets and brain, 65kD)	M81882	1.41
fibromodulin	NM_002023	1.41
homeo box D9	NM_014213	1.41
Consensus includes gb:BF111268 /FEA=EST /DB_XREF=gi:10940958 /DB_XREF=est:7174d02.x1 /CLONE=IMAGE:3527090 /UG=Hs.12436 Homo sapiens cDNA: FLJ22656 fis, clone HSI076	BF111268	1.41
polyadenylate binding protein-interacting protein 1	BF248165	1.41
VENT-like homeobox 2	NM_014468	1.41
troponin T2, cardiac	X79857	1.41
suppressor of variegation 3-9 (Drosophila) homolog 2; hypothetical protein FLJ23414	NM_024670	1.41
hypothetical protein FLJ23604	NM_025064	1.41
intersectin 1 (SH3 domain protein)	NM_003024	1.41
rhodopsin kinase	NM_002929	1.41
Consensus includes gb:AK024909.1 /DEF=Homo sapiens cDNA: FLJ21256 fis, clone COL01402. /FEA=mRNA /DB_XREF=gi:10437324 /UG=Hs.306724 Homo sapiens cDNA: FLJ21256 fis, cl	AK024909	1.41
tenascin XB	M25813	1.41
immune dysregulation, polyendocrinopathy, enteropathy, X-linked	NM_014009	1.41
secreted phosphoprotein 1 (osteopontin, bone sialoprotein I, early T-lymphocyte activation 1)	M83248	1.41
Consensus includes gb:AK021418.1 /DEF=Homo sapiens cDNA FLJ11356 fis, clone HEMBA1000150, highly similar to Homo sapiens putative RNA helicase mRNA. /FEA=mRNA /DB_XREF=gi	AK021418	1.41
TNTSP for transmembrane molecule with thrombospondin module	NM_018676	1.41
fibroblast growth factor 23	NM_020638	1.41
pancreatic beta cell growth factor	NM_012277	1.41
cyclin D2	AI635187	1.41
SRY (sex determining region Y)-box 4	BG528420	1.41
dipeptidylpeptidase IV (CD26, adenosine deaminase complexing protein 2)	M74777	1.41
hypothetical protein FLJ10058	NM_017985	1.41
Consensus includes gb:AK025276.1 /DEF=Homo sapiens cDNA: FLJ21623 fis, clone COL07915. /FEA=mRNA /DB_XREF=gi:10437759 /UG=Hs.306791 Homo sapiens cDNA: FLJ21623 fis, cl	AK025276	1.41
RAB35, member RAS oncogene family	NM_006861	1.41
RNA binding motif, single stranded interacting protein 3	NM_014483	1.41
Consensus includes gb:AW452357 /FEA=EST /DB_XREF=gi:6993133 /DB_XREF=est:UI-H-BI3-alr-d-03-0-UI.s1 /CLONE=IMAGE:3068213 /UG=Hs.27373 Homo sapiens mRNA; cDNA DKFZp5	AW452357	1.41
Itchy homolog E3 ubiquitin protein ligase (mouse)	AB056663	1.41
potassium inwardly-rectifying channel,subfamily J, member 13	AB013889	1.41
Slab-interacting protein	BC005975	1.41
eukaryotic translation termination factor 1	NM_004730	1.41
Consensus includes gb:AF070575.1 /DEF=Homo sapiens clone 24407 mRNA sequence. /FEA=mRNA /DB_XREF=gi:3387946 /UG=Hs.12432 Homo sapiens clone 24407 mRNA sequence	AF070575	1.41
LIM homeobox transcription factor 1, beta	NM_002316	1.41
KIAA0551 protein	AF172268	1.41
Consensus includes gb:AK026793.1 /DEF=Homo sapiens cDNA: FLJ23140 fis, clone LNG09065. /FEA=mRNA /DB_XREF=gi:10439729 /UG=Hs.287728 Homo sapiens cDNA: FLJ23140 fis, cl	AK026793	1.40
collagen, type XIV, alpha 1 (undulin)	M64108	1.40
hypothetical protein FLJ10097	AL523320	1.40
Consensus includes gb:AA156777 /FEA=EST /DB_XREF=gi:1728392 /DB_XREF=est:zd18c08.s1 /CLONE=IMAGE:502286 /UG=Hs.25431 KIAA1219 protein	AA156777	1.40

GTP-binding protein Rho7	BF511718	1.40
KIAA0800 gene product	NM_014703	1.40
angiopoietin-like factor	NM_021146	1.40
gb:NM_022125.1 /DEF=Homo sapiens immortalization-upregulated protein (IMUP), mRNA. /FEA=mRNA /GEN=IMUP /PROD=immortalization-upregulated protein /DB_XREF=gi:11545848 /(	NM_022125	1.40
DKFZP564P1916 protein	NM_015652	1.40
hypothetical protein FLJ23590	NM_024649	1.40
protein kinase, AMP-activated, beta 2 non-catalytic subunit	NM_005399	1.40
syntaphilin	NM_014723	1.40
glycogenin 2	U94364	1.40
interleukin 17 (cytotoxic T-lymphocyte-associated serine esterase 8)	Z58820	1.40
CGI-142	AL133102	1.40
interleukin 1 receptor accessory protein-like 1	NM_014271	1.40
synaptojanin 1	AB020717	1.40
Consensus includes gb:AA029811 /FEA=EST /DB_XREF=gi:1496048 /DB_XREF=est:zk07e10.s1 /CLONE=IMAGE:469866 /UG=Hs.79531 Homo sapiens TTF-I interacting peptide 20 mRNA,	AA029811	1.40
tryptase beta 2	NM_024164	1.40
CD8 antigen, alpha polypeptide (p32)	AW006735	1.40
phospholipase A2, group V	NM_000929	1.40
killer cell immunoglobulin-like receptor, two domains, long cytoplasmic tail, 2	L76669	1.40
gamma-aminobutyric acid (GABA) A receptor, delta	NM_000815	1.40
potassium voltage-gated channel, shaker-related subfamily, beta member 3	NM_004732	1.40
small proline-rich protein 1B (cornifin)	NM_003125	1.40
hypothetical protein FLJ23049	NM_024687	1.40
Consensus includes gb:AK024369.1 /DEF=Homo sapiens cDNA FLJ14307 fis, clone PLACE3000158. /FEA=mRNA /DB_XREF=gi:10436739 /UG=Hs.181341 Homo sapiens cDNA FLJ14307 fi:	AK024369	1.40
nucleosomal binding protein 1	NM_030763	1.40
hypothetical protein MGC8407	NM_024046	1.40
hypothetical protein PRO1163	NM_018576	1.40
keratin associated protein 9.9	NM_030975	1.40
KDEL (Lys-Asp-Glu-Leu) endoplasmic reticulum protein retention receptor 3	NM_016657	1.40
heat shock transcription factor 2 binding protein	NM_007031	1.40
hypothetical protein	NM_019021	1.40
Consensus includes gb:AL133386 /DEF=Human DNA sequence from clone RP1-181C24 on chromosome 6p11.1-12.2. Contains the 3 end of the BMP5 gene for bone morphogenetic protein	AL133386	1.40
tumor protein 63 kDa with strong homology to p53	AF091627	1.40
KIAA0316 gene product	AW297087	1.40
ryanodine receptor 1 (skeletal)	NM_000540	1.40
H factor 1 (complement)	X04697	1.40
KIAA1324 protein	AB037745	1.40
hypothetical protein FLJ12078	NM_024977	1.40
Consensus includes gb:AL080134.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434G043 (from clone DKFZp434G043). /FEA=mRNA /DB_XREF=gi:5262575 /UG=Hs.168762 Homo sapiens mI	AL080134	1.40
fasciculation and elongation protein zeta 1 (zygin I)	NM_022549	1.40
thioredoxin reductase beta	AB019695	1.40
gb:NM_003902.1 /DEF=Homo sapiens far upstream element (FUSE) binding protein 1 (FUBP1), mRNA. /FEA=mRNA /GEN=FUBP1 /PROD=far upstream element-binding protein /DB_XREF:	NM_003902	1.40
ribonuclease, RNase A family, 1 (pancreatic)	NM_002933	1.40
Consensus includes gb:X55989 /DEF=Human ECRP gene for eosinophil cationic related protein /FEA=mRNA /DB_XREF=gi:22940 /UG=Hs.166037 Human ECRP gene for eosinophil cationic	X55989	1.40
Consensus includes gb:AI186145 /FEA=EST /DB_XREF=gi:3736783 /DB_XREF=est:qe46a06.x1 /CLONE=IMAGE:1742002 /UG=Hs.154336 Homo sapiens clone 24583 mRNA sequence	AI186145	1.40
putative opioid receptor, neuromedin K (neurokinin B) receptor-like	NM_006679	1.40
regulator of G-protein signalling 3	NM_021106	1.40
Consensus includes gb:AL137435.1 /DEF=Homo sapiens mRNA; cDNA DKFZp761G0924 (from clone DKFZp761G0924). /FEA=mRNA /DB_XREF=gi:6807994 /UG=Hs.274577 Homo sapiens	AL137435	1.40
small nuclear RNA activating complex, polypeptide 1, 43kD	NM_003082	1.40
hypothetical protein FLJ22056	NM_022489	1.40
ectonucleoside triphosphate diphosphohydrolase 2	NM_001246	1.40
peroxisomal biogenesis factor 3	AU157140	1.40
metal-regulatory transcription factor 1	AW182367	1.40
DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 3	NM_001356	1.40
HDCMA18P protein	AK000089	1.40
hypothetical protein PRTD-NY3	AY009107	1.40
iduronate 2-sulfatase (Hunter syndrome)	AF179281	1.40
f-box and leucine-rich repeat protein 6	NM_024555	1.40
neuregulin 1	L12260	1.40
KIAA0981 protein	AB023198	1.40
Consensus includes gb:AK024615.1 /DEF=Homo sapiens cDNA: FLJ20962 fis, clone ADSH00804. /FEA=mRNA /DB_XREF=gi:10436931 /UG=Hs.306698 Homo sapiens cDNA: FLJ20962 fis,	AK024615	1.40
Consensus includes gb:AL121900 /DEF=Human DNA sequence from clone RP11-379J5 on chromosome 20 Contains the last exon of the SEC23B gene for Sec23 (S. cerevisiae) homolog B, a	AL121900	1.40
pregnancy specific beta-1-glycoprotein 9	M94890	1.40
gene amplified in squamous cell carcinoma 1	AI341811	1.40

transglutaminase 5	NM_004245	1.40
nuclear receptor interacting protein 1	NM_003489	1.40
Consensus includes gb:AU150691 /FEA=EST /DB_XREF=gi:11012212 /DB_XREF=est:AU150691 /CLONE=NT2RP2003367 /UG=Hs.153321 Homo sapiens cDNA FLJ10577 fis, clone NT2RP2	AU150691	1.40
Consensus includes gb:AI417917 /FEA=EST /DB_XREF=gi:4261421 /DB_XREF=est:tg55d04.x1 /CLONE=IMAGE:2112679 /UG=Hs.20733 Homo sapiens cDNA: FLJ22356 fis, clone HRC0634	AI417917	1.40
Consensus includes gb:AK025564.1 /DEF=Homo sapiens cDNA: FLJ21911 fis, clone HEP03855. /FEA=mRNA /DB_XREF=gi:10438116 /UG=Hs.306816 Homo sapiens cDNA: FLJ21911 fis, cl	AK025564	1.40
gb:BC005043.1 /DEF=Homo sapiens, Similar to tumor necrosis factor receptor superfamily, member 10c, decoy without an intracellular domain, clone MGC:12787, mRNA, complete cds. /FE	BC005043	1.40
protein C (inactivator of coagulation factors Va and VIIIa)	NM_000312	1.40
KIAA0993 protein	AI806395	1.40
prothymosin, alpha (gene sequence 28)	I04798	1.40
gb:BC000233.1 /DEF=Homo sapiens, Similar to guanine nucleotide binding protein, alpha transducing 2, clone MGC:2232, mRNA, complete cds. /FEA=mRNA /PROD=Similar to guanine nucl	BC000233	1.40
T cell receptor gamma locus	M30894	1.40
Consensus includes gb:AK024890.1 /DEF=Homo sapiens cDNA: FLJ21237 fis, clone COL01114. /FEA=mRNA /DB_XREF=gi:10437303 /UG=Hs.306720 Homo sapiens cDNA: FLJ21237 fis, cl	AK024890	1.40
secreted protein, acidic, cysteine-rich (osteonectin)	AL575922	1.40
interleukin 9 receptor	Z84723	1.40
gamma-aminobutyric acid (GABA) A receptor, alpha 1	NM_000806	1.40
type I intermediate filament cytokeratin	NM_015515	1.40
ATP synthase, H+ transporting, mitochondrial F0 complex, subunit e	NM_007100	1.40
angiotensin I converting enzyme (peptidyl-dipeptidase A) 2	AK026461	1.40
cyclin G associated kinase	D88435	1.40
hypothetical protein FLJ14103	NM_024689	1.40
LUC7-like (S. cerevisiae)	NM_018032	1.40
unactive progesterone receptor, 23 kD	BE903880	1.40
collagen, type I, alpha 1	Y15916	1.40
protein phosphatase 1, regulatory (inhibitor) subunit 14D	NM_017726	1.40
desmocollin 3	NM_001941	1.40
keratin 2A (epidermal ichthyosis bullosa of Siemens)	NM_000423	1.40
MAD, mothers against decapentaplegic homolog 7 (Drosophila)	NM_005904	1.40
cytochrome P450, subfamily XIX (aromatization of androgens)	NM_000103	1.40
protein phosphatase 3 (formerly 2B), catalytic subunit, gamma isoform (calcineurin A gamma)	BE670265	1.40
BAI1-associated protein 2	NM_017451	1.40
glycophorin B (includes Ss blood group)	U05255	1.40
dead ringer-like 1 (Drosophila)	NM_005224	1.40
Consensus includes gb:W87634 /FEA=EST /DB_XREF=gi:1401698 /DB_XREF=est:zh67b09.s1 /CLONE=IMAGE:417113 /UG=Hs.17947 ESTs, Weakly similar to K02F3.10 C.elegans	W87634	1.40
KIAA0882 protein	AI348094	1.40
leukocyte receptor cluster (LRC) member 4	NM_024298	1.40
hypothetical protein MGC5487	BC001305	1.40
kallikrein 2, prostatic	AF188747	1.40
betaine-homocysteine methyltransferase	NM_001713	1.40
ATP-binding cassette, sub-family A (ABC1), member 4	U88667	1.40
carcinoembryonic antigen-related cell adhesion molecule 1 (biliary glycoprotein)	M76742	1.40
hypothetical protein FLJ20401	NM_017805	1.40
junctophilin 3	NM_020655	1.40
spectrin, beta, non-erythrocytic 1	NM_003128	1.40
teratocarcinoma-derived growth factor 1	NM_003212	1.40
KIAA1042 protein	AK000754	1.40
deleted in azoospermia	AF271088	1.40
Consensus includes gb:BF000251 /FEA=EST /DB_XREF=gi:10700526 /DB_XREF=est:7h24e02.x1 /CLONE=IMAGE:3316922 /UG=Hs.227835 KIAA1049 protein	BF000251	1.40
oculocerebrorenal syndrome of Lowe	NM_000276	1.40
Consensus includes gb:AL137378.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434K1126 (from clone DKFZp434K1126). /FEA=mRNA /DB_XREF=gi:6807908 /UG=Hs.306455 Homo sapiens	AL137378	1.40
v-abl Abelson murine leukemia viral oncogene homolog 2 (arg, Abelson-related gene)	NM_007314	1.40
adenosine A3 receptor	NM_000677	1.40
seizure related 6 homolog (mouse)-like	NM_021115	1.40
CDC14 cell division cycle 14 homolog B (S. cerevisiae)	AF064105	1.40
NAD(P)H dehydrogenase, quinone 1	NM_000903	1.40
cytochrome P450, subfamily IIC (mephenytoin 4-hydroxylase), polypeptide 9	AV657878	1.40
KIAA1609 protein	AA195017	1.40
malic enzyme 1, NADP(+)-dependent, cytosolic	NM_002395	1.40
retinoic acid induced 17	AF070622	1.40
Tax interaction protein 1	AK001327	1.40
hypothetical protein FLJ20013	AK026097	1.40
RAB31, member RAS oncogene family	AF183421	1.40
NK3 transcription factor homolog A (Drosophila)	AF249671	1.40
Consensus includes gb:AK025352.1 /DEF=Homo sapiens cDNA: FLJ21699 fis, clone COL09829. /FEA=mRNA /DB_XREF=gi:10437851 /UG=Hs.306804 Homo sapiens cDNA: FLJ21699 fis, cl	AK025352	1.40

bromodomain containing protein 7S kDa human homolog	NM_012107	1.40
TAF4A RNA polymerase II, TATA box binding protein (TBP)-associated factor, 135 kD	AI744029	1.40
Consensus includes gb:AK026856.1 /DEF=Homo sapiens cDNA: FLJ23203 fis, clone ADKA02487. /FEA=mRNA /DB_XREF=gi:10439812 /UG=Hs.306889 Homo sapiens cDNA: FLJ23203 fis,	AK026856	1.40
serum-inducible kinase	NM_006622	1.40
KIAA0626 gene product	AF327560	1.40
zinc finger protein 291	AW139369	1.40
KIAA1046 protein	T79216	1.40
potassium large conductance calcium-activated channel, subfamily M, beta member 2	NM_005832	1.40
testes-specific protease 50	NM_013270	1.40
hypothetical protein FLJ14251	AA504646	1.40
phosphoglycerate kinase 1	AL121974	1.40
Consensus includes gb:AL442084.1 /DEF=Homo sapiens mRNA; cDNA DKFZp547G044 (from clone DKFZp547G044). /FEA=mRNA /DB_XREF=gi:10241763 /UG=Hs.306924 Homo sapiens n	AL442084	1.40
Consensus includes gb:AA160522 /FEA=EST /DB_XREF=gi:1735889 /DB_XREF=est:zq46b10.s1 /CLONE=IMAGE:632731 /UG=Hs.301667 Homo sapiens mRNA; cDNA DKFZp566I043 (from	AA160522	1.40
lipopolysaccharide binding protein	NM_004139	1.40
phospholamban	NM_002667	1.40
nebulette	NM_006393	1.40
basic leucine zipper transcription factor, ATF-like	NM_006399	1.40
alcohol dehydrogenase 6 (class V)	NM_000672	1.40
trans-golgi network protein 2	NM_006464	1.40
solute carrier family 4, sodium bicarbonate cotransporter, member 4	AF157492	1.40
keratin, cuticle, ultrahigh sulphur 1	NM_005553	1.40
histamine receptor H3	NM_007232	1.40
angiopoietin 1	NM_001146	1.40
hypothetical protein MGC2889	BC005856	1.40
hypothetical protein FLJ10803	AA554945	1.40
hypothetical protein FLJ11292	AK023539	1.40
Consensus includes gb:Z93929 /DEF=Human DNA sequence from clone 272E8 on chromosome Xp22.13-22.31. Contains a pseudogene similar to MDM2-Like P53-binding protein gene. Cont	Z93929	1.40
mitogen-activated protein kinase kinase kinase 9	AF251442	1.40
mitochondrial ribosomal protein S11	AL109953	1.40
dual specificity phosphatase 3 (vaccinia virus phosphatase VH1-related)	AL048503	1.40
potassium intermediate/small conductance calcium-activated channel, subfamily N, member 3	AI251016	1.40
phospholipid scramblase 3	NM_020360	1.40
ribonuclease, RNase A family, 3 (eosinophil cationic protein)	NM_002935	1.40
mitogen-activated protein kinase 10	NM_002753	1.40
KIAA0669 gene product	AF201292	1.40
sirtuin silent mating type information regulation 2 homolog 3 (S. cerevisiae)	AI492888	1.40
Consensus includes gb:AK024128.1 /DEF=Homo sapiens cDNA FLJ14066 fis, clone HEMBB1001197. /FEA=mRNA /DB_XREF=gi:10436433 /UG=Hs.306665 Homo sapiens cDNA FLJ14066 f	AK024128	1.40
cleavage and polyadenylation specific factor 1, 160kD subunit	BE676642	1.40
potassium large conductance calcium-activated channel, subfamily M, beta member 4	NM_014505	1.40
hypothetical protein FLJ22714	NM_025050	1.40
G protein-coupled receptor 21	NM_005294	1.40
lymphocyte antigen 94 homolog, activating NK-receptor; NK-p46, (mouse)	NM_004829	1.39
killer cell immunoglobulin-like receptor, two domains, short cytoplasmic tail, 1	NM_014512	1.39
Lutheran blood group (Auberger b antigen included)	X83425	1.39
Carbonic anhydrase-related protein 10	NM_020178	1.39
gamma-aminobutyric acid (GABA) A receptor, beta 1	NM_000812	1.39
hypothetical protein FLJ20126	NM_004686	1.39
endoplasmic reticulum resident protein 58; hypothetical protein MGC5302	NM_024089	1.39
major histocompatibility complex, class I, F	BE138825	1.39
putative chemokine receptor; GTP-binding protein	NM_006018	1.39
epithelial protein lost in neoplasm beta	NM_016357	1.39
CCR4-NOT transcription complex, subunit 7	NM_013354	1.39
KIAA0322 protein	AB048365	1.39
major histocompatibility complex, class II, DQ alpha 1	X00452	1.39
Consensus includes gb:AL122074.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434N0435 (from clone DKFZp434N0435); partial cds. /FEA=mRNA /GEN=DKFZp434N0435 /PROD=hypothetic	AL122074	1.39
connexin-36	NM_020660	1.39
TU3A protein	NM_007177	1.39
schwannomin interacting protein 1	NM_014575	1.39
Consensus includes gb:AK023058.1 /DEF=Homo sapiens cDNA FLJ12996 fis, clone NT2RP3000235. /FEA=mRNA /DB_XREF=gi:10434801 /UG=Hs.287546 Homo sapiens cDNA FLJ12996 fi	AK023058	1.39
cerberus 1 homolog, cysteine knot superfamily (Xenopus laevis)	NM_005454	1.39
retinoic acid induced 2	NM_021785	1.39
myotubularin related protein 7	AF073482	1.39
potassium inwardly-rectifying channel, subfamily J, member 2	AF153820	1.39

Consensus includes gb:AF070571.1 /DEF=Homo sapiens clone 24739 mRNA sequence. /FEA=mRNA /DB_XREF=gi:3387940 /UG=Hs.44690 Homo sapiens clone 24739 mRNA sequence	AF070571	1.39
KIAA0711 gene product	NM_014867	1.39
HSPC072 protein	NM_014162	1.39
collagen, type V, alpha 1	N30339	1.39
ATP-binding cassette, sub-family A (ABC1), member 3	NM_001089	1.39
brain expressed, X-linked 1	NM_018476	1.39
hypothetical protein FLJ20581	NM_017888	1.39
ryanodine receptor 2 (cardiac)	BE968750	1.39
Rho guanine exchange factor (GEF) 15	NM_014958	1.39
Rho guanine nucleotide exchange factor (GEF) 5	NM_005435	1.39
hypothetical protein FLJ23231	NM_025079	1.39
peptidylprolyl isomerase C (cyclophilin C)	NM_000943	1.39
chitinase 3-like 1 (cartilage glycoprotein-39)	M80927	1.39
gb:U19348.1 /DEF=Human (tpr-met fusion) oncogene mRNA, complete cds. /FEA=mRNA /GEN=tprmet fusion /PROD=tpr-met fusion protein /DB_XREF=gi:625085 /FL=gb:U19348.1	U19348	1.39
F-box and WD-40 domain protein 7 (archipelago homolog, Drosophila)	NM_018315	1.39
PP1201 protein	NM_022152	1.39
Down syndrome critical region gene 1	NM_004414	1.39
Consensus includes gb:AL096729.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434D044 (from clone DKFZp434D044). /FEA=mRNA /DB_XREF=gi:5419863 /UG=Hs.146581 Homo sapiens mI	AL096729	1.39
Consensus includes gb:BF967657 /FEA=EST /DB_XREF=gi:12334872 /DB_XREF=est:602287358T1 /CLONE=IMAGE:4374495 /UG=Hs.90005 superiorcervical ganglia, neural specific 10 /FI	BF967657	1.39
lymphocyte antigen 117	AI735692	1.39
zinc finger protein 294	AL163248	1.39
Rhesus blood group, CcEe antigens	X63095	1.39
Consensus includes gb:AL049381.1 /DEF=Homo sapiens mRNA; cDNA DKFZp586I2118 (from clone DKFZp586I2118). /FEA=mRNA /DB_XREF=gi:4500168 /UG=Hs.21851 Homo sapiens cD	AL049381	1.39
hippocalcin-like protein 4	AL136591	1.39
interleukin 6 signal transducer (gp130, oncostatin M receptor)	AB015706	1.39
chorionic somatomammotropin hormone 1 (placental lactogen)	NM_022640	1.39
homeo box A5	NM_019102	1.39
olfactory receptor, family 1, subfamily D, member 4	NM_003552	1.39
4-hydroxyphenylpyruvate dioxygenase	NM_002150	1.39
CCR4-NOT transcription complex, subunit 4	AF180475	1.39
tumor necrosis factor (ligand) superfamily, member 13	BF448647	1.39
solute carrier family 7 (cationic amino acid transporter, y+ system), member 9	NM_014270	1.39
TEK tyrosine kinase, endothelial (venous malformations, multiple cutaneous and mucosal)	NM_000459	1.39
hypothetical protein FLJ20581	AI733019	1.39
ceroid-lipofuscinosis, neuronal 3, juvenile (Batten, Spielmeyer-Vogt disease)	AC002544	1.39
C-type lectin-like receptor-1	NM_016511	1.39
glutathione S-transferase M3 (brain)	AL527430	1.39
atrophin-1 interacting protein 1; activin receptor interacting protein 1	NM_012301	1.39
ribosomal protein S6 kinase, 70kD, polypeptide 2	NM_003952	1.39
KIAA1109 protein	AL137254	1.39
bromodomain adjacent to zinc finger domain, 2A	NM_013449	1.39
low density lipoprotein receptor (familial hypercholesterolemia)	S70123	1.39
Consensus includes gb:BC002892.1 /DEF=Homo sapiens, clone IMAGE:3940152, mRNA, partial cds. /FEA=mRNA /PROD=Unknown (protein for IMAGE:3940152) /DB_XREF=gi:12804078 /L	BC002892	1.39
Consensus includes gb:AI002715 /FEA=EST /DB_XREF=gi:3203129 /DB_XREF=est:an20d12.s1 /CLONE=IMAGE:1699223 /UG=Hs.130315 ESTs	AI002715	1.39
protocadherin beta 3	NM_018937	1.39
hypothetical protein	AI005473	1.39
serine/threonine kinase 38	AF034187	1.39
angiotensin receptor-like 2	NM_005162	1.39
hypothetical protein FLJ20701	NM_017933	1.39
keratin 12 (Meesmann corneal dystrophy)	NM_000223	1.39
Consensus includes gb:BF434846 /FEA=EST /DB_XREF=gi:11447134 /DB_XREF=est:7o74d07.x1 /CLONE=IMAGE:3641917 /UG=Hs.296757 Homo sapiens cDNA: FLJ20933 fis, clone ADSE	BF434846	1.39
HEAT-like repeat-containing protein	NM_020958	1.39
protease, serine, 3 (trypsin 3)	NM_002771	1.39
phosphatidic acid phosphatase type 2A	AB000888	1.39
immunoglobulin lambda locus	X79782	1.39
hypothetical protein FLJ20069	AL136797	1.39
Consensus includes gb:AF035314.1 /DEF=Homo sapiens clone 23651 mRNA sequence. /FEA=mRNA /DB_XREF=gi:2661076 /UG=Hs.134526 Homo sapiens clone 23651 mRNA sequence	AF035314	1.39
transcription factor 3 (E2A immunoglobulin enhancer binding factors E12/E47)	AL117663	1.39
prostate differentiation factor	BC000529	1.39
KIAA1547 protein	AI567426	1.39
Consensus includes gb:AA558468 /FEA=EST /DB_XREF=gi:2328945 /DB_XREF=est:nk39c06.s1 /CLONE=IMAGE:1015882 /UG=Hs.178665 Homo sapiens immunoglobulin lambda gene loci	AA558468	1.39
uromodulin (uromucoid, Tamm-Horsfall glycoprotein)	NM_003361	1.39
homogentisate 1,2-dioxygenase (homogentisate oxidase)	NM_000187	1.39

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IL-17B receptor	NM_018725	1.39
factor H-related protein 5	NM_030787	1.39
G protein-coupled receptor 52	NM_005684	1.39
sentrin/SUMO-specific protease 3	AK000923	1.39
wingless-type MMTV integration site family, member 2B	NM_024494	1.39
prostate stem cell antigen	NM_005672	1.39
epilepsy, progressive myoclonus type 2, Lafora disease (laforin)	NM_005670	1.39
myeloid/lymphoid or mixed-lineage leukemia (trithorax homolog, Drosophila)	AA704766	1.39
gb:NM_016425.1 /DEF=Homo sapiens transmembrane protease, serine 4 (TMPRSS4), mRNA. /FEA=mRNA /GEN=TMPRSS4 /PROD=transmembrane protease, serine 4 /DB_XREF=gi:77059;	NM_016425	1.39
Consensus includes gb:AA400421 /FEA=EST /DB_XREF=gi:2054292 /DB_XREF=est:zu69h09.r1 /CLONE=IMAGE:743297 /UG=Hs.55896 Homo sapiens PAC clone RP5-978E18 from 7p21	AA400421	1.39
vasoactive intestinal peptide receptor 2	X95097	1.39
peptidylglycine alpha-amidating monooxygenase	BF038548	1.39
neuromedin U	NM_006681	1.39
histamine H4 receptor	NM_021624	1.39
Consensus includes gb:AK022999.1 /DEF=Homo sapiens cDNA FLJ12937 fis, clone NT2RP2005020. /FEA=mRNA /DB_XREF=gi:10434712 /UG=Hs.14846 Homo sapiens mRNA; cDNA DKFZ	AK022999	1.39
bromodomain adjacent to zinc finger domain, 2A	BE513659	1.39
zinc finger protein 42 (myeloid-specific retinoic acid- responsive)	AI733248	1.39
MTA22 protein	BF031714	1.39
Consensus includes gb:AL049265.1 /DEF=Homo sapiens mRNA; cDNA DKFZp564F053 (from clone DKFZp564F053). /FEA=mRNA /DB_XREF=gi:4500013 /UG=Hs.71968 Homo sapiens mR	AL049265	1.39
T cell receptor alpha locus	AE000659	1.39
Consensus includes gb:BE794663 /FEA=EST /DB_XREF=gi:10215861 /DB_XREF=est:601590459F1 /CLONE=IMAGE:3944463 /UG=Hs.306668 Homo sapiens cDNA FLJ14089 fis, clone MAI	BE794663	1.39
KIAA0546 protein	AB011118	1.39
proline-rich protein BstNI subfamily 3	NM_006249	1.39
putative GR6 protein	NM_007354	1.39
calcitonin gene-related peptide-receptor component protein	NM_014478	1.39
KIAA1157 protein	AB032983	1.39
hypothetical protein FLJ20489	AW149696	1.39
syntrophin, beta 2 (dystrophin-associated protein A1, 59kD, basic component 2)	AI762021	1.39
insulin-like growth factor binding protein 4	NM_001552	1.39
Cdc42 effector protein 2	BC005406	1.39
hypothetical protein FLJ22222	NM_024648	1.39
APG5 autophagy 5-like (S. cerevisiae)	NM_004849	1.39
matrix metalloproteinase 12 (macrophage elastase)	NM_002426	1.39
ATP-binding cassette, sub-family G (WHITE), member 1	U34919	1.39
LIM domain binding 3	AA211481	1.39
syndecan 1	NM_002997	1.39
transducer of ERBB2, 2	D64109	1.39
tryptase, alpha	AF206665	1.39
DKFZP564I122 protein	BC006122	1.39
fucosyltransferase 6 (alpha (1,3) fucosyltransferase)	U27336	1.39
zinc finger protein 200	AU149447	1.39
Consensus includes gb:AK025650.1 /DEF=Homo sapiens cDNA: FLJ21997 fis, clone HEP06590. /FEA=mRNA /DB_XREF=gi:10438234 /UG=Hs.306820 Homo sapiens cDNA: FLJ21997 fis, cl	AK025650	1.39
ribosomal protein L5	AA582460	1.39
TAF1 RNA polymerase II, TATA box binding protein (TBP)-associated factor, 250 kD	X07024	1.39
proprotein convertase subtilisin/kexin type 5	NM_006200	1.39
SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily d, member 3	NM_003078	1.39
nidogen (enactin)	NM_002508	1.39
hypothetical protein FLJ20706	NM_017935	1.39
Consensus includes gb:BF046750 /FEA=EST /DB_XREF=gi:10763806 /DB_XREF=est:INSSA4 /CLONE=227760 /UG=Hs.274454 ESTs, Weakly similar to 2004399A chromosomal protein H.s	BF046750	1.39
N-acetylated alpha-linked acidic dipeptidase-like; ILEAL DIPEPTIDYLPEPTIDASE	NM_005468	1.39
nuclear receptor subfamily 0, group B, member 1	NM_000475	1.39
tryptase beta 1	AF206667	1.39
mutS homolog 5 (E. coli)	BE674960	1.39
hypothetical protein DKFZp434P144	AL136868	1.39
Consensus includes gb:AA837026 /FEA=EST /DB_XREF=gi:2912225 /DB_XREF=est:og09c12.s1 /CLONE=IMAGE:1439350 /UG=Hs.163549 ESTs	AA837026	1.39
Fc fragment of IgG, high affinity Ia, receptor for (CD64)	X14355	1.39
ribosomal protein S6 kinase, 90kD, polypeptide 5	AF074393	1.39
poly (ADP-ribose) glycohydrolase	NM_003631	1.39
cyfidin, basic protein of sperm head cytoskeleton 2	NM_001340	1.39
hypothetical protein FLJ12606	NM_024804	1.39
H2A histone family, member I	NM_003511	1.39
butyrylcholinesterase	NM_000055	1.39
olfactory receptor, family 3, subfamily A, member 1	NM_002550	1.39

hypothetical protein FLJ13114	NM_024541	1.39
KIAA0792 gene product	NM_014698	1.39
KIAA0367 protein	AL138349	1.39
small inducible cytokine subfamily A (Cys-Cys), member 24	NM_002991	1.38
calpain 9 (nCL-4)	NM_006615	1.38
hemoglobin, alpha 2	V00489	1.38
hypothetical protein DKFZp434K1172	NM_031290	1.38
proteasome (prosome, macropain) subunit, alpha type, 7	NM_002792	1.38
immunoglobulin heavy constant mu	X95660	1.38
KIAA0758 protein	BF941499	1.38
eukaryotic translation initiation factor 2, subunit 1 (alpha, 35kD)	AA577698	1.38
lectin, galactoside-binding, soluble, 8 (galectin 8)	AF342816	1.38
KIAA0513 gene product	NM_014732	1.38
Consensus includes gb:U85992.1 /DEF=Human clone IMAGE:35527 unknown protein mRNA, partial cds. /FEA=mRNA /PROD=unknown /DB_XREF=gi:1835745 /UG=Hs.87197 Human clone	U85992	1.38
KIAA0215 gene product	NM_014735	1.38
insulin promoter factor 1, homeodomain transcription factor	U35632	1.38
KIAA0574 protein	AB011146	1.38
protein tyrosine kinase 9	N25562	1.38
aryl hydrocarbon receptor nuclear translocator	NM_001668	1.38
Consensus includes gb:R25849 /FEA=EST /DB_XREF=gi:781984 /DB_XREF=est:yh42a11.r1 /CLONE=IMAGE:132380 /UG=Hs.217305 ESTs	R25849	1.38
fatty-acid-Coenzyme A ligase, long-chain 3	NM_004457	1.38
ephrin-B3	U57001	1.38
kallikrein 10	BC002710	1.38
parvin, beta	AA187563	1.38
CD86 antigen (CD28 antigen ligand 2, B7-2 antigen)	NM_006889	1.38
a disintegrin and metalloproteinase domain 21	NM_003813	1.38
ALL1-fused gene from chromosome 1q	BC006471	1.38
KIAA0461 protein	AK000199	1.38
putative dimethyladenosine transferase	NM_014473	1.38
chromosome 8 open reading frame 1	AI738702	1.38
T cell receptor beta locus	U66059	1.38
hypothetical protein PRO1489	NM_018584	1.38
hypothetical protein IMPACT	NM_018439	1.38
nuclear receptor subfamily 4, group A, member 1	D85245	1.38
hypothetical protein FLJ22558	NM_003858	1.38
interferon, omega 1	NM_002177	1.38
hypothetical protein FLJ20584	NM_017891	1.38
secreted frizzled-related protein 1	NM_003012	1.38
Consensus includes gb:L42230 /DEF=Homo sapiens core 2 beta-1,6-N-acetylglucosaminyltransferase pseudogene /FEA=CDS /DB_XREF=gi:1019163 /UG=Hs.247990 Homo sapiens core 2	L42230	1.38
Consensus includes gb:AF218021.1 /DEF=Homo sapiens clone PP552 unknown mRNA. /FEA=mRNA /PROD=unknown /DB_XREF=gi:10441971 /UG=Hs.302127 Homo sapiens clone PP552	AF218021	1.38
KIAA1655 protein	AB051442	1.38
hypothetical protein FLJ23074	NM_025052	1.38
zinc finger protein 281	NM_012482	1.38
insulin-like 5	NM_005478	1.38
chromodomain helicase DNA binding protein 3	NM_001272	1.38
integrin, beta 6	NM_000888	1.38
ephrin-A5	NM_001962	1.38
Consensus includes gb:AL137285.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434D2416 (from clone DKFZp434D2416). /FEA=mRNA /DB_XREF=gi:6807740 /UG=Hs.306445 Homo sapiens	AL137285	1.38
Consensus includes gb:AK024142.1 /DEF=Homo sapiens cDNA FLJ14080 fis, clone HEMBB1002152. /FEA=mRNA /DB_XREF=gi:10436451 /UG=Hs.301905 Homo sapiens cDNA FLJ14080 f	AK024142	1.38
natural killer cell transcript 4	NM_004221	1.38
hypothetical protein MGC14258	AV717623	1.38
thioredoxin reductase 2	AI247566	1.38
heme oxygenase (decycling) 1	NM_002133	1.38
Consensus includes gb:AL137798 /DEF=Human DNA sequence from clone RP5-1182A14 on chromosome 1 Contains part of a gene similar to rat Espin, a pseudogene similar to KIAA0454, a	AL137798	1.38
ATPase, Class VI, type 11A	BF439472	1.38
lymphocyte cytosolic protein 2 (SH2 domain-containing leukocyte protein of 76kD)	NM_005565	1.38
ret finger protein-like 3 antisense	AL021937	1.38
KIAA0638 protein	AB014538	1.38
p53-induced protein	NM_006034	1.38
nuclear receptor coactivator 3	AF010227	1.38
integral membrane protein 2A	NM_004867	1.38
son of sevenless homolog 1 (Drosophila)	L13857	1.38
Jun dimerization protein p21SNFT	NM_018664	1.38

calmodulin 1 (phosphorylase kinase, delta)	NM_006888	1.38
keratin, hair, acidic,2	NM_002278	1.38
pyruvate dehydrogenase kinase, isoenzyme 3	NM_005391	1.38
synaptojanin 2	AK026758	1.38
flavin containing monooxygenase 5	NM_001461	1.38
hypothetical protein FLJ11168	NM_018347	1.38
CD58 antigen, (lymphocyte function-associated antigen 3)	D28586	1.38
KIAA0632 protein	AK001615	1.38
poliovirus receptor-related 1 (herpesvirus entry mediator C; nectin)	AF110314	1.38
adipose specific 2	NM_006829	1.38
interleukin 26	NM_018402	1.38
myotubularin related protein 3	NM_021090	1.38
PTD011 protein	NM_014051	1.38
neuropilin 2	AF280546	1.38
guanylate cyclase 1, soluble, beta 3	AF020340	1.38
HC6/V-6 protein	AF036973	1.38
collagen, type VI, alpha 3	NM_004369	1.38
glutamate receptor, ionotropic, N-methyl D-aspartate 2C	L76224	1.38
Consensus includes gb:BF508244 /FEA=EST /DB_XREF=gi:11591542 /DB_XREF=est:UI-H-BI4-aqa-b-02-0-UI.s1 /CLONE=IMAGE:3089210 /UG=Hs.300832 ESTs, Highly similar to I53872 d	BF508244	1.38
retinoblastoma-like 2 (p130)	X76061	1.38
hypothetical protein FLJ20972	NM_025030	1.38
synaptotagmin I	AV731490	1.38
immunoglobulin lambda locus	D01059	1.38
gap junction protein, alpha 7, 45kD (connexin 45)	NM_005497	1.38
KIAA0419 gene product	NM_014711	1.38
Consensus includes gb:AK024945.1 /DEF=Homo sapiens cDNA: FLJ21292 fis, clone COL01969. /FEA=mRNA /DB_XREF=gi:10437367 /UG=Hs.306732 Homo sapiens cDNA: FLJ21292 fis, cl	AK024945	1.38
a disintegrin and metalloproteinase domain 12 (meltrin alpha)	NM_021641	1.38
Consensus includes gb:AB051440.1 /DEF=Homo sapiens mRNA for KIAA1653 protein, partial cds. /FEA=mRNA /GEN=KIAA1653 /PROD=KIAA1653 protein /DB_XREF=gi:13359178 /UG=Hs.	AB051440	1.38
KIAA0841 protein	AW189966	1.38
claudin 6	NM_021195	1.38
defensin, alpha 6, Paneth cell-specific	NM_001926	1.38
Consensus includes gb:AL121903 /DEF=Human DNA sequence from clone RP1-155G6 on chromosome 20 Contains part of the gene for brefeldin A-inhibited guanine nucleotide-exchange pr	AL121903	1.38
H factor (complement)-like 3	NM_005666	1.38
Consensus includes gb:AL133089.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434E2028 (from clone DKFZp434E2028). /FEA=mRNA /DB_XREF=gi:6453541 /UG=Hs.274562 Homo sapiens	AL133089	1.38
estrogen receptor 1	M12674	1.38
hypothetical protein FLJ10547	NM_018134	1.38
DKFZP564G2022 protein	BG493972	1.38
TAL1 (SCL) interrupting locus	NM_003035	1.38
serum/glucocorticoid regulated kinase 2	NM_016276	1.38
complement component 1, q subcomponent, alpha polypeptide	NM_015991	1.38
spastic paraplegia 4 (autosomal dominant; spastin)	NM_014946	1.38
alcohol dehydrogenase 1B (class I), beta polypeptide	AF153821	1.38
advillin	AJ796813	1.38
CD58 antigen, (lymphocyte function-associated antigen 3)	NM_001779	1.38
tyrosinase (oculocutaneous albinism IA)	NM_000372	1.38
caspase 1, apoptosis-related cysteine protease (interleukin 1, beta, convertase)	U13699	1.38
epidermal growth factor receptor pathway substrate 15	NM_001981	1.38
v-myc myelocytomatosis viral oncogene homolog 1, lung carcinoma derived (avian)	M19720	1.38
desmocollin 2	BF196457	1.38
mannose-binding lectin (protein A) 1, pseudogene 1	AF019382	1.38
alpha-2-glycoprotein 1, zinc	AC004522	1.38
gb:D26121.1 /DEF=Human mRNA for ZFM1 protein alternatively spliced product, complete cds. /FEA=mRNA /PROD=ZFM1 protein, alternatively spliced product /DB_XREF=gi:785998 /UG=	D26121	1.38
Consensus includes gb:AI907083 /FEA=EST /DB_XREF=gi:6497611 /DB_XREF=est:PM-BT134-050499-650 /UG=Hs.124620 ESTs	AI907083	1.38
intracellular hyaluronan-binding protein	AF241831	1.38
mucin 4, tracheobronchial	AJ242547	1.38
potassium voltage-gated channel, delayed-rectifier, subfamily S, member 1	NM_002251	1.38
defensin, beta 1	U73945	1.38
ubiquitin specific protease 1	AW499935	1.38
hypothetical protein FLJ10916	NM_018271	1.38
hypothetical protein FLJ22028	NM_024854	1.38
annexin A3	M63310	1.38
SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily a, member 4	AA772023	1.38
4-rmatopontin	AJ146848	1.38

acid sphingomyelinase-like phosphodiesterase	NM_014474	1.38
S164 protein	AI925305	1.38
complement component 7	NM_000587	1.38
proteolipid protein 1 (Pelizaeus-Merzbacher disease, spastic paraplegia 2, uncomplicated)	BC002665	1.38
Consensus includes gb:AL109726.1 /DEF=Homo sapiens mRNA full length insert cDNA clone EUROIMAGE 124430. /FEA=mRNA /DB_XREF=gi:5689792 /UG=Hs.187513 Homo sapiens mRN	AL109726	1.38
C-reactive protein, pentraxin-related	NM_000567	1.38
runt-related transcription factor 2	NM_004348	1.38
gb:NM_018628.1 /DEF=Homo sapiens hypothetical protein PRO2472 (PRO2472), mRNA. /FEA=mRNA /GEN=PRO2472 /PROD=hypothetical protein PRO2472 /DB_XREF=gi:8924169 /UG=	NM_018628	1.38
nuclear antigen Sp100	AF056322	1.38
CD36 antigen (collagen type I receptor, thrombospondin receptor)	NM_000072	1.38
crystallin, gamma A	NM_014617	1.38
dom-3 homolog 2 (C. elegans)	NM_005510	1.38
hypothetical protein MGC16714	AL117502	1.38
phospholipase A2, group IID	NM_012400	1.38
serine (or cysteine) proteinase inhibitor, clade C (antithrombin), member 1	D29832	1.38
amyloid beta precursor protein (cytoplasmic tail) binding protein 2	AA046411	1.38
hypothetical protein FLJ20736	NM_017948	1.38
PRO1748 protein	AF118073	1.38
adenovirus 5 E1A binding protein	BE250417	1.38
tumor necrosis factor receptor superfamily, member 4	NM_003327	1.38
calpain 6	NM_014289	1.38
zinc finger protein 147 (estrogen-responsive finger protein)	NM_005082	1.38
hypothetical protein FLJ10038	NM_017976	1.38
solute carrier family 23 (nucleobase transporters), member 1	AF164142	1.38
transient receptor potential channel 5	NM_012471	1.38
acetylserotonin O-methyltransferase	NM_004043	1.38
hypothetical protein FLJ14146	NM_024709	1.38
periplakin	NM_002705	1.38
acyl-Coenzyme A oxidase 1, palmitoyl	S69189	1.38
transmembrane protein vezatin	NM_017599	1.38
PYK2 N-terminal domain-interacting receptor 1	NM_031220	1.38
taste receptor, type 2, member 13	NM_023920	1.38
hypothetical protein FLJ14345	NM_024733	1.38
protein tyrosine phosphatase, receptor type, J	AI806482	1.38
ubiquitin specific protease 2	AF079564	1.38
hypothetical protein FLJ12377	NM_024989	1.38
heat shock 70kD protein 1A	NM_005345	1.38
pregnancy specific beta-1-glycoprotein 6	M31125	1.38
hypothetical protein FLJ11556	NM_024964	1.38
Consensus includes gb:AL535113 /FEA=EST /DB_XREF=gi:12798606 /DB_XREF=est:AL535113 /CLONE=CS0DF008YC23 (3 prime) /UG=Hs.283006 phospholipase C, beta 4 /FL=gb:NM_0	AL535113	1.38
aquaporin 4	D63412	1.38
cytochrome P450, subfamily IIIA, polypeptide 7	AF315325	1.38
nuclear factor of activated T-cells 5, tonicity-responsive	AI005683	1.38
RAB27A, member RAS oncogene family	U38654	1.38
G protein-coupled receptor 37 (endothelin receptor type B-like)	U87460	1.38
tumor necrosis factor (ligand) superfamily, member 10	NM_003810	1.38
glutamate decarboxylase 1 (brain, 67kD)	NM_000817	1.38
KIAA0560 gene product	AB011132	1.38
potassium inwardly-rectifying channel, subfamily J, member 8	BF514158	1.38
KIAA0779 protein	AB018322	1.38
acyloxyacyl hydrolase (neutrophil)	NM_001637	1.38
oncostatin M receptor	NM_003999	1.38
hypothetical protein FLJ10871	NM_018250	1.38
nuclear factor I/C (CCAAT-binding transcription factor)	X12492	1.38
retinol binding protein 1, cellular	NM_002899	1.38
signal transduction protein (SH3 containing)	NM_005864	1.38
Consensus includes gb:AI185160 /FEA=EST /DB_XREF=gi:3735798 /DB_XREF=est:qe30g10.s1 /CLONE=IMAGE:1740546 /UG=Hs.326248 Homo sapiens cDNA: FLJ22071 fis, clone HEP11	AI185160	1.38
transcription factor 4	NM_003199	1.38
cyclic AMP-regulated phosphoprotein, 21 kD	NM_016300	1.38
tumor endothelial marker 8	NM_018153	1.38
interleukin 12B (natural killer cell stimulatory factor 2, cytotoxic lymphocyte maturation factor 2, p40)	NM_002187	1.38
hypothetical protein FLJ22659	NM_024934	1.38
uncharacterized hypothalamus protein HSMNP1	NM_018478	1.38

Consensus includes gb:U38460 /DEF=Human RNA binding motif (RBM) gene, partial cds /FEA=CDS /DB_XREF=gi:1061399 /UG=Hs.326802 Human RNA binding motif (RBM) gene, partial c	U38460	1.38
EGF-containing fibulin-like extracellular matrix protein 2	NM_016938	1.38
TATA box binding protein (TBP)-associated factor, RNA polymerase I, B, 63kD	AK026521	1.38
gb:Z22814.1 /DEF=H.sapiens ORF, complete CDS. /FEA=mRNA /DB_XREF=gi:393697 /FL=gb:Z22814.1	Z22814	1.38
KIAA1548 protein	NM_020909	1.38
hypothetical protein FLJ10803	NM_018224	1.38
myelin basic protein	M13577	1.38
M24537B subtilis pheB, phoA genes corresponding to nucleotides 2017-3334 of M24537 (-5, -M, -3 represent transcript regions 5 prime, Middle, and 3 prime respectively)	M24537B	1.38
Huntingtin interacting protein H	AF161412	1.38
casein kinase 1, gamma 2	NM_001319	1.38
cold autoinflammatory syndrome 1	AK027194	1.37
T-cell leukemia/lymphoma 6	NM_020552	1.37
PTPL1-associated RhoGAP 1	NM_004815	1.37
hypothetical protein FLJ11808	NM_024728	1.37
Consensus includes gb:AK025247.1 /DEF=Homo sapiens cDNA: FLJ21594 fis, clone COL07053. /FEA=mRNA /DB_XREF=gi:10437718 /UG=Hs.288571 Homo sapiens cDNA: FLJ21594 fis, cl	AK025247	1.37
agouti signaling protein, nonagouti homolog (mouse)	NM_001672	1.37
RAD54-like (S.cerevisiae)	NM_003579	1.37
extracellular link domain-containing 1	AL574194	1.37
solute carrier family 12 (sodium/chloride transporters), member 3	AI627943	1.37
Rho GTPase activating protein 1	U62794	1.37
Homer, neuronal immediate early gene, 1B	BE550452	1.37
KIAA1659 protein	AB051446	1.37
cysteine-rich motor neuron 1	NM_016441	1.37
biliverdin reductase A	NM_000712	1.37
mitogen-activated protein kinase 11	NM_002751	1.37
KIAA0373 gene product	NM_014684	1.37
nuclear autoantigen	AF243424	1.37
gb:BC006367.1 /DEF=Homo sapiens, clone MGC:12878, mRNA, complete cds. /FEA=mRNA /PROD=Unknown (protein for MGC:12878) /DB_XREF=gi:13623518 /FL=gb:BC006367.1	BC006367	1.37
calcium/calmodulin-dependent protein kinase IG	AL049688	1.37
Consensus includes gb:AI927479 /FEA=EST /DB_XREF=gi:5663443 /DB_XREF=est:wo90d06.x1 /CLONE=IMAGE:2462603 /UG=Hs.165590 ribosomal protein S13	AI927479	1.37
retinoic acid receptor, alpha	AI806984	1.37
sine oculis homeobox homolog 2 (Drosophila)	NM_016932	1.37
TU3A protein	AL050264	1.37
gb:U36759.1 /DEF=Human pre-T cell receptor alpha-type chain precursor, mRNA, complete cds. /FEA=mRNA /PROD=pre-T cell receptor alpha-type chain precursor /DB_XREF=gi:1127580	U36759	1.37
Consensus includes gb:BF229395 /FEA=EST /DB_XREF=gi:11166291 /DB_XREF=est:MR1-CI0019-260900-001-a11 /UG=Hs.159454 ESTs	BF229395	1.37
fructosamine-3-kinase	NM_022158	1.37
translocase of inner mitochondrial membrane 17 homolog A (yeast)	NM_006335	1.37
microtubule-associated protein 7	T62571	1.37
formin-binding protein 3	Z78308	1.37
integrin, alpha V (vitronectin receptor, alpha polypeptide, antigen CD51)	AI093579	1.37
potassium inwardly-rectifying channel, subfamily J, member 4	U24056	1.37
cysteine knot superfamily 1, BMP antagonist 1	AF154054	1.37
eukaryotic translation initiation factor 2C, 1	NM_012199	1.37
zinc transporter	NM_014579	1.37
fatty acid binding protein 4, adipocyte	NM_001442	1.37
Consensus includes gb:Y14488.1 /DEF=Homo sapiens mRNA for putative 14kD protein containing SHMT homology, clone pUS1215. /FEA=mRNA /PROD=14 kDa protein /DB_XREF=gi:583(	Y14488	1.37
HSPC054 protein	NM_014152	1.37
trinucleotide repeat containing 3	NM_005878	1.37
transmembrane 7 superfamily member 1 (upregulated in kidney)	NM_003272	1.37
ATP synthase, H+ transporting, mitochondrial F1 complex, epsilon subunit	NM_006886	1.37
natural killer-tumor recognition sequence	AI361805	1.37
synaptosomal-associated protein, 23kD	NM_003825	1.37
Consensus includes gb:AL162044.1 /DEF=Homo sapiens mRNA; cDNA DKFZp761L0812 (from clone DKFZp761L0812); partial cds. /FEA=mRNA /GEN=DKFZp761L0812 /PROD=hypothetica	AL162044	1.37
Consensus includes gb:BF222018 /FEA=EST /DB_XREF=gi:11129195 /DB_XREF=est:7p41a04.x1 /CLONE=IMAGE:3648271 /UG=Hs.289803 ESTs, Moderately similar to S71105 protein-gl	BF222018	1.37
KIAA0426 gene product	Z98745	1.37
hypothetical protein MGC2628	NM_024076	1.37
FOS-like antigen-1	BG251266	1.37
T-box 1	NM_005992	1.37
KIAA0527 protein	BF977837	1.37
KIAA0446 gene product	AL526243	1.37
malic enzyme 3, NADP(+)-dependent, mitochondrial	NM_006680	1.37
GABA(A) receptors associated protein like 3	AF180519	1.37
cholinergic receptor, nicotinic, alpha polypeptide 4	X89745	1.37

phosphoglucomutase 5	NM_021965	1.37
formyl peptide receptor-like 1	M88107	1.37
putative leukocyte platelet-activating factor receptor	NM_022571	1.37
hypothetical protein FLJ21120	NM_024767	1.37
hypothetical protein MGC4294	NM_024314	1.37
mal, T-cell differentiation protein	NM_002371	1.37
hypothetical protein FLJ10851	NM_018245	1.37
KIAA0451 gene product	NM_014826	1.37
KIAA0015 gene product	NM_014634	1.37
Consensus includes gb:AW675655 /FEA=EST /DB_XREF=gi:7540890 /DB_XREF=est:ba52e01.x1 /CLONE=IMAGE:2900184 /UG=Hs.314158 ESTs	AW675655	1.37
hypothetical protein MGC13138	AA746290	1.37
gb:NM_025111.1 /DEF=Homo sapiens hypothetical protein FLJ23571 (FLJ23571), mRNA. /FEA=mRNA /GEN=FLJ23571 /PROD=hypothetical protein FLJ23571 /DB_XREF=gi:13376682 /UG	NM_025111	1.37
KIAA0186 gene product	NM_021067	1.37
endothelin converting enzyme 2	NM_014693	1.37
Bacillus subtilis /REF=L38424 /DEF=B subtilis dapB, jojF, jojG genes corresponding to nucleotides 2634-3089 of L38424 /LEN=1931 (-5, -M, -3 represent transcript regions 5 prime, Midd	L38424	1.37
purinergic receptor P2Y, G-protein coupled, 1	NM_002563	1.37
Consensus includes gb:BF447272 /FEA=EST /DB_XREF=gi:11512410 /DB_XREF=est:7p47a12.x1 /CLONE=IMAGE:3648695 /UG=Hs.288190 ESTs	BF447272	1.37
solute carrier family 6 (neurotransmitter transporter, betaine/GABA), member 12	U27699	1.37
Fc fragment of IgE, low affinity II, receptor for (CD23A)	NM_002002	1.37
organic cation transporter	NM_020372	1.37
spleen focus forming virus (SFFV) proviral integration oncogene spi1	NM_003120	1.37
Consensus includes gb:AW168915 /FEA=EST /DB_XREF=gi:6400440 /DB_XREF=est:xj15b10.x1 /CLONE=IMAGE:2657275 /UG=Hs.283946 Homo sapiens prostate-specific membrane antig	AW168915	1.37
NA618 protein	AF210651	1.37
C21orf19-like protein	NM_015955	1.37
hypothetical protein FLJ12810	NM_022836	1.37
hypothetical protein MGC4090	NM_024310	1.37
forkhead box O3A	AF041336	1.37
hypothetical protein FLJ11598	NM_024966	1.37
calcium channel, voltage-dependent, alpha 1I subunit	NM_021096	1.37
Consensus includes gb:AK024949.1 /DEF=Homo sapiens cDNA: FLJ21296 fis, clone COL02029. /FEA=mRNA /DB_XREF=gi:10437372 /UG=Hs.306734 Homo sapiens cDNA: FLJ21296 fis, cl	AK024949	1.37
adaptor-related protein complex 4, epsilon 1 subunit	NM_007347	1.37
caspase recruitment domain protein 14	NM_024110	1.37
decorin	AI281593	1.37
trans-golgi network protein 2	BF061845	1.37
DKFZP586N0721 protein	NM_015400	1.37
PRO0159 protein	NM_014118	1.37
suppressor of Ty 3 homolog (S. cerevisiae)	AF064804	1.37
hypothetical protein FLJ13511	AF209931	1.37
POU domain, class 4, transcription factor 2	NM_004575	1.37
collagen, type XIII, alpha 1	M59217	1.37
G protein-coupled receptor 12	NM_005288	1.37
solute carrier family 16 (monocarboxylic acid transporters), member 6	NM_004694	1.37
hypothetical protein FLJ10829	NM_018234	1.37
G protein-coupled receptor	AI817041	1.37
islet cell autoantigen 1 (69kD)	L21181	1.37
zinc finger protein 180 (HHZ168)	NM_013256	1.37
phosphodiesterase 6G, cGMP-specific, rod, gamma	M36476	1.37
ADP-ribosylation factor-like 3	NM_004311	1.37
testis specific leucine rich repeat protein	NM_012472	1.37
activin A receptor, type I	NM_001105	1.37
H1 histone family, member T (testis-specific)	NM_005323	1.37
sulfotransferase family, cytosolic, 2A, dehydroepiandrosterone (DHEA) -preferring, member 1	NM_003167	1.37
transthyretin (prealbumin, amyloidosis type I)	AF162690	1.37
DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 24	BG421209	1.37
hemopexin	M36803	1.37
heat shock transcription factor 2	M65217	1.37
Consensus includes gb:BG109555 /FEA=EST /DB_XREF=gi:12603061 /DB_XREF=est:602280658F1 /CLONE=IMAGE:4367966 /UG=Hs.331207 ESTs	BG109555	1.37
sodium channel, nonvoltage-gated 1, delta	NM_002978	1.37
wingless-type MMTV integration site family, member 5A	NM_003392	1.37
syntaxin 6	NM_005819	1.37
olfactory receptor, family 10, subfamily H, member 2	NM_013939	1.37
amyloid beta (A4) precursor protein-binding, family A, member 2 (X11-like)	AB014719	1.37
retinol binding protein 4, plasma	AA908769	1.37

olfactory receptor, family 2, subfamily C, member 1	NM_012368	1.37
Fc fragment of IgG, low affinity IIb, receptor for (CD32)	U90941	1.37
Wilms tumor associated protein	NM_015855	1.37
CGI-60 protein	NM_016008	1.37
DnaJ (Hsp40) homolog, subfamily B, member 12	AI695173	1.37
sema domain, immunoglobulin domain (Ig), transmembrane domain (TM) and short cytoplasmic domain, (semaphorin) 4F	NM_004263	1.37
KIAA0712 gene product	BC000277	1.37
feline sarcoma oncogene	NM_002005	1.37
ureidopropionase, beta	NM_016327	1.37
fatty acid binding protein 6, ileal (gastrotropin)	U19869	1.37
hypothetical protein FLJ12988	NM_024999	1.37
Consensus includes gb:BF223370 /FEA=EST /DB_XREF=gi:11130547 /DB_XREF=est:7q87e03.x1 /CLONE=IMAGE:3705316 /UG=Hs.90449 Human clone 23908 mRNA sequence	BF223370	1.37
KIAA0878 protein	AK023621	1.37
contactin associated protein-like 2	AU144598	1.37
microfibrillar-associated protein 2	AL049569	1.37
neuropeptide Y receptor Y5	NM_006174	1.37
X04603 B subtilis thrC, thrB genes corresponding to nucleotides 248-2229 of X04603 (-5, -M, -3 represent transcript regions 5 prime, Middle, and 3 prime respectively)	X04603	1.37
phospholipid scramblase 1	AI825926	1.37
Consensus includes gb:AF007147.1 /DEF=Homo sapiens clone 23712 mRNA sequence. /FEA=mRNA /DB_XREF=gi:2852625 /UG=Hs.113166 Homo sapiens clone 23712 mRNA sequence	AF007147	1.37
claudin 18	NM_016369	1.37
hypothetical protein DKFZp547M136 similar to widely-interspaced zinc finger motifs	AI089655	1.37
Consensus includes gb:AW973253 /FEA=EST /DB_XREF=gi:8163100 /DB_XREF=est:EST385351 /UG=Hs.292689 ESTs	AW973253	1.37
lymphocyte antigen 9	AL582804	1.37
inter-alpha (globulin) inhibitor H4 (plasma Kallikrein-sensitive glycoprotein)	NM_002218	1.37
palmdelphin	NM_017734	1.37
Consensus includes gb:AW242315 /FEA=EST /DB_XREF=gi:6576069 /DB_XREF=est:xm96d12.x1 /CLONE=IMAGE:2692055 /UG=Hs.27860 Homo sapiens mRNA; cDNA DKFZp586M0723 (fr	AW242315	1.37
Consensus includes gb:AK021571.1 /DEF=Homo sapiens cDNA FLJ11509 fis, clone HEMBA1002166. /FEA=mRNA /DB_XREF=gi:10432777 /UG=Hs.289093 Homo sapiens cDNA FLJ11509 fi	AK021571	1.37
echinoderm microtubule-associated protein-like	NM_004434	1.37
hematopoietic PBX-interacting protein	BF344265	1.37
B lymphoid tyrosine kinase	BC004473	1.37
hypothetical protein FLJ12681	AL515916	1.37
lymphotoxin beta receptor (TNFR superfamily, member 3)	NM_002342	1.37
EphB2	AF025304	1.37
Consensus includes gb:AK025271.1 /DEF=Homo sapiens cDNA: FLJ21618 fis, clone COL07487. /FEA=mRNA /DB_XREF=gi:10437753 /UG=Hs.306790 Homo sapiens cDNA: FLJ21618 fis, cl	AK025271	1.37
myosin VI	NM_004999	1.37
complement component 5	NM_001735	1.37
short-chain alcohol dehydrogenase family member	NM_005794	1.37
CGI-146 protein	AK025651	1.37
transporter protein; system N1 Na+ and H+-coupled glutamine transporter	NM_006841	1.37
DKFZP564F1422 protein	AL049942	1.37
testis-specific transcript, Y-linked 2	AF000991	1.37
mitogen-activated protein kinase 6	NM_002748	1.37
sema domain, immunoglobulin domain (Ig), short basic domain, secreted, (semaphorin) 3B	NM_004636	1.37
myosin, heavy polypeptide 8, skeletal muscle, perinatal	M36769	1.37
contactin associated protein-like 2	AC005378	1.37
NGFI-A binding protein 2 (EGR1 binding protein 2)	BF337329	1.37
KIAA0831 protein	NM_014924	1.37
hypothetical protein DKFZp434F0318	NM_030817	1.37
hypothetical protein FLJ22833	NM_022837	1.37
KIAA1096 protein	AI359472	1.37
HMG-box transcription factor TCF-3	NM_031283	1.37
fucosyltransferase 6 (alpha (1,3) fucosyltransferase)	U27331	1.37
cyclin-dependent kinase inhibitor 2C (p18, inhibits CDK4)	U17074	1.37
3'-phosphoadenosine 5'-phosphosulfate synthase 2	NM_004670	1.37
KIAA0878 protein	NM_014899	1.37
SH3-domain binding protein 1	NM_018957	1.37
chondroitin 4-sulfotransferase	NM_018413	1.37
carbonic anhydrase XII	NM_001218	1.37
p53-regulated apoptosis-inducing protein 1	NM_022112	1.37
leukocyte immunoglobulin-like receptor, subfamily A (without TM domain), member 5	NM_024317	1.37
DKFZP586N0819 protein	AI689429	1.37
Norrie disease (pseudoglioma)	NM_000266	1.37
Consensus includes gb:AI378979 /FEA=EST /DB_XREF=gi:4188832 /DB_XREF=est:tc40d07.x1 /CLONE=IMAGE:2067085 /UG=Hs.292911 ESTs	AI378979	1.37

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ATP-binding cassette, sub-family C (CFTR/MRP), member 3	AK000791	1.37
activated p21cdc42Hs kinase	AK024904	1.37
xanthene dehydrogenase	U06117	1.37
Consensus includes gb:R73554 /FEA=EST /DB_XREF=gi:847586 /DB_XREF=est:yj94e05.s1 /CLONE=IMAGE:156416 /UG=Hs.103391 Human insulin-like growth factor binding protein 5 (IG	R73554	1.37
FOX2 forkhead factor	NM_018416	1.37
sperm adhesion molecule 1 (PH-20 hyaluronidase, zona pellucida binding)	L13779	1.37
regulator of G-protein signalling 5	AF159570	1.37
tumor necrosis factor receptor superfamily, member 10b	BC001281	1.37
BMX non-receptor tyrosine kinase	NM_001721	1.37
hypothetical protein FLJ13055	AA209463	1.36
SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily a, member 4	AI831675	1.36
Consensus includes gb:AA700015 /FEA=EST /DB_XREF=gi:2702978 /DB_XREF=est:zi69c11.s1 /CLONE=IMAGE:436052 /UG=Hs.287995 Homo sapiens cDNA: FLJ23181 fis, clone LNG1109	AA700015	1.36
testis-specific kinase substrate	NM_021733	1.36
coronin, actin binding protein, 2A	NM_003389	1.36
CEGP1 protein	AI424243	1.36
plexin A2	NM_025179	1.36
tripartite motif-containing 29	NM_012101	1.36
fibroblast growth factor 2 (basic)	M27968	1.36
mitochondrial capsule selenoprotein	NM_030663	1.36
putative protein-tyrosine kinase	NM_015978	1.36
WAP four-disulfide core domain 2	NM_006103	1.36
ret finger protein-like 3 antisense	AL021937	1.36
small proline-rich protein 2B	NM_006945	1.36
U2 small nuclear ribonucleoprotein auxiliary factor (65kD)	NM_007279	1.36
carbohydrate (N-acetylglucosamine 6-O) sulfotransferase 7	AL022165	1.36
alpha-methylacyl-CoA racemase	AF047020	1.36
protein kinase C, mu	NM_002742	1.36
aminolevulinate, delta-, synthase 2 (sideroblastic/hypochromic anemia)	Z83821	1.36
6 protein-coupled receptor C5L2	NM_018485	1.36
hypothetical protein FLJ23510	NM_024720	1.36
DKFZP586P0123 protein	AL080220	1.36
hypothetical protein FLJ22814	NM_024916	1.36
protein kinase C and casein kinase substrate in neurons 3	NM_016223	1.36
Cluster Incl. AI801973:tz29d05.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-2270985 /clone_end=3 /gb=AI801973 /gi=5367445 /ug=Hs.27373 /len=532	AI801973	1.36
KIAA0563 gene product	NM_014834	1.36
nuclear factor of activated T-cells, cytoplasmic, calcineurin-dependent 3	U85430	1.36
S100 calcium binding protein A5	NM_002962	1.36
v-yes-1 Yamaguchi sarcoma viral oncogene homolog 1	NM_005433	1.36
unr-interacting protein	NM_007178	1.36
calpain 3, (p94)	AF127764	1.36
hypothetical protein DKFZp434M1923	NM_030974	1.36
major histocompatibility complex, class II, DP alpha 1	AI128225	1.36
singlepass-type MMTV integration site family, member 6	AI191771	1.36
Consensus includes gb:AF339787.1 /DEF=Homo sapiens clone IMAGE:205688, mRNA sequence. /FEA=mRNA /DB_XREF=gi:13507325 /UG=Hs.326722 Homo sapiens clone IMAGE:205688,	AF339787	1.36
Consensus includes gb:AW503390 /FEA=EST /DB_XREF=gi:7118730 /DB_XREF=est:UI-HF-BN0-akw-c-02-0-UI.r1 /CLONE=IMAGE:3078507 /UG=Hs.135258 ESTs, Weakly similar to ALU7_H	AW503390	1.36
BCL2-like 11 (apoptosis facilitator)	NM_006538	1.36
interleukin 1 receptor, type II	NM_004633	1.36
chromodomain helicase DNA binding protein 2	NM_001271	1.36
cholinergic receptor, nicotinic, delta polypeptide	NM_000751	1.36
interleukin 24	NM_006850	1.36
KIAA1058 protein	AW450751	1.36
UDP-GlcNAc:betaGal beta-1,3-N-acetylglucosaminyltransferase 3	NM_014256	1.36
nephrin related gene	NM_018240	1.36
Consensus includes gb:AF203728.1 /DEF=Homo sapiens bone morphogenetic protein BMPY (BMPY) mRNA, partial cds. /FEA=mRNA /GEN=BMPY /PROD=bone morphogenetic protein BMP'	AF203728	1.36
collagen, type XI, alpha 1	I04177	1.36
gamma-aminobutyric acid (GABA) A receptor, gamma 2	NM_000816	1.36
calcium binding protein 1 (calbrain)	AF169148	1.36
ephrin-B2	BF001670	1.36
Gardner-Rasheed feline sarcoma viral (v-fgr) oncogene homolog	BC002836	1.36
NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 7 (18kD, B18)	M33374	1.36
solute carrier family 21 (organic anion transporter), member 3	NM_021094	1.36
epoxide hydrolase 1, microsomal (xenobiotic)	NM_000120	1.36
vesicle trafficking, beach and anchor containing	AI659561	1.36

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sialyltransferase	NM_006456	1.36
LIS1-interacting protein NUDE1, rat homolog	NM_017668	1.36
methyl CpG binding protein 2 (Rett syndrome)	NM_004992	1.36
KIAA0669 gene product	AF201292	1.36
gamma-aminobutyric acid (GABA) A receptor, alpha 4	NM_000809	1.36
PRO0255 protein	NM_014124	1.36
carbonic anhydrase XIV	NM_012113	1.36
cerebellin 1 precursor	NM_004352	1.36
interleukin 15	NM_000585	1.36
Consensus includes gb:AK024879.1 /DEF=Homo sapiens cDNA: FLJ21226 fis, clone COL00721. /FEA=mRNA /DB_XREF=gi:10437291 /UG=Hs.306715 Homo sapiens cDNA: FLJ21226 fis, cl	AK024879	1.36
tryptase delta 1	NM_012217	1.36
TU1281-TY protein	NM_016575	1.36
Consensus includes gb:AL080215.1 /DEF=Homo sapiens mRNA; cDNA DKFZp586I0323 (from clone DKFZp586I0323). /FEA=mRNA /DB_XREF=gi:5262706 /UG=Hs.102301 Homo sapiens n	AL080215	1.36
myelin transcription factor 1-like	AL133024	1.36
B-cell CLL/lymphoma 11A (zinc finger protein)	NM_022893	1.36
5-hydroxytryptamine (serotonin) receptor 1B	M81590	1.36
centrosomal protein 2	BC001433	1.36
interleukin 1 receptor, type I	NM_000877	1.36
growth differentiation factor 9	NM_005260	1.36
chromosome 22 open reading frame 1	NM_001585	1.36
BAI1-associated protein 1	NM_004742	1.36
KIAA0337 gene product	NM_014786	1.36
putative N-acetyltransferase Camello 2	NM_016347	1.36
mitogen-activated protein kinase kinase 4	AA810268	1.36
alkylation repair; alkB homolog	NM_006020	1.36
cytochrome P450, subfamily IIB (phenobarbital-inducible), polypeptide 7	M29873	1.36
a disintegrin and metalloproteinase domain 30	NM_021794	1.36
Consensus includes gb:AL577024 /FEA=EST /DB_XREF=gi:12939748 /DB_XREF=est:AL577024 /CLONE=CS0DI082YN19 (3 prime) /UG=Hs.7921 Homo sapiens mRNA; cDNA DKFZp566E18	AL577024	1.36
hypothetical protein MGC5528	L11372	1.36
retinaldehyde binding protein 1	NM_000326	1.36
transcobalamin I (vitamin B12 binding protein, R binder family)	NM_001062	1.36
lactate dehydrogenase A -like	AY009108	1.36
tumor protein D52	AA524023	1.36
ectodysplasin 1, anhidrotic receptor	NM_022336	1.36
killer cell lectin-like receptor subfamily B, member 1	NM_002258	1.36
trefoil factor 3 (intestinal)	NM_003226	1.36
opioid receptor, delta 1	NM_000911	1.36
hypothetical protein FLJ10508	NM_018118	1.36
ubiquitously transcribed tetratricopeptide repeat gene, Y chromosome	AF000995	1.36
3'-phosphoadenosine 5'-phosphosulfate synthase 2	AW299958	1.36
mucin 1, transmembrane	AF348143	1.36
matrix metalloproteinase 23A	NM_004659	1.36
dual-specificity tyrosine-(Y)-phosphorylation regulated kinase 1A	U52373	1.36
aldo-keto reductase family 1, member D1 (delta 4-3-ketosteroid-5-beta-reductase)	NM_005989	1.36
B-factor, properdin	AF349679	1.36
Wolf-Hirschhorn syndrome candidate 1-like 1	NM_023034	1.36
period homolog 2 (Drosophila)	NM_022817	1.36
inhibitor of growth family, member 1	AW193656	1.36
Kv channel-interacting protein 1	NM_014592	1.36
solute carrier family 12 (potassium/chloride transporters), member 6	NM_005135	1.36
lymphocyte antigen 94 homolog, activating NK-receptor, NK-p46, (mouse)	AJ006121	1.36
nudix (nucleoside diphosphate linked moiety X)-type motif 3	AF062530	1.36
olfactory receptor, family 1, subfamily E, member 1	NM_003553	1.36
L1 cell adhesion molecule (hydrocephalus, stenosis of aqueduct of Sylvius 1, MASA (mental retardation, aphasia, shuffling gait and adducted thumbs) syndrome, spastic paraplegia 1)	NM_000425	1.36
Consensus includes gb:AF003737.1 /DEF=Homo sapiens chromosome 21q22.1 anonymous mRNA sequence. /FEA=mRNA /DB_XREF=gi:2197086 /UG=Hs.302179 Homo sapiens chromoso	AF003737	1.36
matrilin 4	NM_003833	1.36
NAD kinase	BE551347	1.36
nuclear antigen Sp100	NM_003113	1.36
glycoprotein 2 (zymogen granule membrane)	NM_001502	1.36
KIAA0752 protein	AF155097	1.36
calmodulin 3 (phosphorylase kinase, delta)	NM_005184	1.36
ADP-ribosyltransferase (NAD+; poly (ADP-ribose) polymerase)-like 1	NM_006437	1.36
uncoupling protein 3 (mitochondrial, proton carrier)	NM_003356	1.36

protein tyrosine phosphatase, receptor type, N polypeptide 2	U65065	1.36
tumor necrosis factor (ligand) superfamily, member 5 (hyper-IgM syndrome)	NM_000074	1.36
Consensus includes gb:AF055018.1 /DEF=Homo sapiens clone 24442 mRNA sequence. /FEA=mRNA /DB_XREF=gi:3005745 /UG=Hs.139137 Homo sapiens clone 24442 mRNA sequence	AF055018	1.36
Consensus includes gb:AV651241 /FEA=EST /DB_XREF=gi:9872255 /DB_XREF=est:AV651241 /CLONE=GLCCNG03 /UG=Hs.282469 ESTs	AV651241	1.36
phosphomannomutase 1	AL023553	1.36
hypothetical protein FLJ22041 similar to FK506 binding proteins	NM_021939	1.36
small inducible cytokine subfamily A (Cys-Cys), member 13	Z77651	1.36
calpain 11	NM_007058	1.36
DNA (cytosine-5-)-methyltransferase 3 alpha	NM_022552	1.36
ankyrin 1, erythrocytic	AI659683	1.36
CD209 antigen-like	AF245219	1.36
KIAA0993 protein	BE348236	1.36
Consensus includes gb:AF052172.1 /DEF=Homo sapiens clone 24617 mRNA sequence. /FEA=mRNA /DB_XREF=gi:3360483 /UG=Hs.64125 Homo sapiens clone 24617 mRNA sequence	AF052172	1.36
hypothetical protein FLJ20452	AL136872	1.36
Consensus includes gb:AK027217.1 /DEF=Homo sapiens cDNA: FLJ23564 fis, clone LNG10773. /FEA=mRNA /DB_XREF=gi:10440296 /UG=Hs.306913 Homo sapiens cDNA: FLJ23564 fis, cl	AK027217	1.36
gamma-glutamyltransferase-like activity 1	NM_004121	1.36
hypothetical protein FLJ10970	NM_018286	1.36
protein kinase C, alpha binding protein	NM_012407	1.36
G protein-coupled receptor kinase-interactor 2	NM_014776	1.36
histamine N-methyltransferase	N40285	1.36
transducin-like enhancer of split 1 (E(sp1) homolog, Drosophila)	NM_005077	1.36
SRY (sex determining region Y)-box 30	NM_007017	1.36
phosphate cytidylyltransferase 1, choline, alpha isoform	NM_005017	1.36
cytochrome P450, subfamily XIA (cholesterol side chain cleavage)	NM_000781	1.36
erythrocyte membrane protein band 4.9 (dematin)	NM_001978	1.36
angiopoietin-like 4	NM_016109	1.36
opioid receptor, mu 1	U12569	1.36
Consensus includes gb:AL136967 /DEF=Human DNA sequence from clone RP1-149M18 on chromosome 6 Contains ESTs, STSs, GSSs and CpG islands. Contains the natural killer cell p44 rela	AL136967	1.36
RAB7, member RAS oncogene family	AK024417	1.36
hypothetical protein FLJ14360	N38751	1.36
Consensus includes gb:AV741657 /FEA=EST /DB_XREF=gi:10859238 /DB_XREF=est:AV741657 /CLONE=CBMALG01 /UG=Hs.154085 leucine zipper protein 1	AV741657	1.36
thymidylate synthetase	BG281679	1.36
nuclear autoantigen	NM_014574	1.36
DKFZP586D0919 protein	N67741	1.36
Consensus includes gb:AK021569.1 /DEF=Homo sapiens cDNA FLJ11507 fis, clone HEMBA1002160. /FEA=mRNA /DB_XREF=gi:10432775 /UG=Hs.314347 Homo sapiens cDNA FLJ11507 fi	AK021569	1.36
KIAA0790 protein	AK025495	1.36
DKFZP434C153 protein	AL080128	1.36
runt-related transcription factor 1 (acute myeloid leukemia 1; aml1 oncogene)	U19601	1.36
solute carrier family 16 (monocarboxylic acid transporters), member 2 (putative transporter)	NM_006517	1.36
natural killer cell receptor, immunoglobulin superfamily member	NM_007053	1.36
speckle-type POZ protein	BF673888	1.36
potassium voltage-gated channel, shaker-related subfamily, member 10	NM_005549	1.36
integrin, beta-like 1 (with EGF-like repeat domains)	NM_004791	1.36
Consensus includes gb:AU144041 /FEA=EST /DB_XREF=gi:11005562 /DB_XREF=est:AU144041 /CLONE=HEMBA1000726 /UG=Hs.293834 Homo sapiens cDNA FLJ11403 fis, clone HEMBA	AU144041	1.36
Huntingtin interacting protein M	AA815276	1.36
Consensus includes gb:AK021690.1 /DEF=Homo sapiens cDNA FLJ11628 fis, clone HEMBA1004238. /FEA=mRNA /DB_XREF=gi:10432926 /UG=Hs.306610 Homo sapiens cDNA FLJ11628 fi	AK021690	1.36
transglutaminase 2 (C polypeptide, protein-glutamine-gamma-glutamyltransferase)	M98478	1.36
mutL homolog 3 (E. coli)	AB039667	1.36
calcium channel, voltage-dependent, beta 3 subunit	U07139	1.36
interleukin 15	Y09908	1.36
immune dysregulation, polyendocrinopathy, enteropathy, X-linked	NM_014009	1.36
Consensus includes gb:BE223030 /FEA=EST /DB_XREF=gi:8910348 /DB_XREF=est:hu46c02.x1 /CLONE=IMAGE:3173090 /UG=Hs.301281 Homo sapiens mRNA, chromosome 1 specific tra	BE223030	1.36
protein kinase, cAMP-dependent, regulatory, type II, alpha	BF246917	1.36
E2F transcription factor 3	AI640363	1.36
killer cell immunoglobulin-like receptor, two domains, long cytoplasmic tail, 4	AF276292	1.36
molybdenum cofactor synthesis 3	NM_014484	1.36
lymphocyte antigen 6 complex, locus D	NM_003695	1.36
guanylate cyclase activator 1A (retina)	L36861	1.36
zinc finger protein 232	NM_014519	1.36
hypothetical protein FLJ20190	NM_017705	1.36
pregnancy specific beta-1-glycoprotein 1	M34715	1.36
KIAA0599 protein	BG223341	1.36
Kruppel-like factor 7 (ubiquitous)	AA488672	1.35

small inducible cytokine subfamily B (Cys-X-Cys), member 10	NM_001565	1.35
POU domain, class 6, transcription factor 1	L14482	1.35
undifferentiated embryonic cell transcription factor 1	NM_003577	1.35
homolog of mouse quaking QKI (KH domain RNA binding protein)	AL031781	1.35
ATPase, H+ transporting, lysosomal (vacuolar proton pump) 42kD	AW024925	1.35
hypothetical protein DKFZp566J091	NM_030915	1.35
KIAA1111 protein	AF091081	1.35
myogenin (myogenic factor 4)	NM_002479	1.35
Consensus includes gb:AL049646 /DEF=Human DNA sequence from clone RP4-568F9 on chromosome 20 Contains the ZNF133 (zinc finger protein 133 (clone pH2-13)) gene, part of a gen	AL049646	1.35
homolog of Xenopus Claspin	NM_022111	1.35
Consensus includes gb:AK024849.1 /DEF=Homo sapiens cDNA: FLJ21196 fis, clone COL00193. /FEA=mRNA /DB_XREF=gi:10437257 /UG=Hs.287650 Homo sapiens cDNA: FLJ21196 fis, cl	AK024849	1.35
Consensus includes gb:AF141901.1 /DEF=Homo sapiens GKAPSAPAP interacting protein (SHANK) mRNA, partial cds. /FEA=mRNA /GEN=SHANK /PROD=GKAPSAPAP interacting protein /DB_	AF141901	1.35
hypothetical protein FLJ10199	NM_018022	1.35
SH3-domain protein 5 (ponsin)	AF356525	1.35
polymyositis/scleroderma autoantigen 1 (75kD)	AI346350	1.35
discoidin domain receptor family, member 1	L11315	1.35
hepatocyte nuclear factor 3, beta	AI693985	1.35
T-box 19	NM_005149	1.35
family with sequence similarity 8, member A1	NM_016255	1.35
transcription factor 7 (T-cell specific, HMG-box)	AW027359	1.35
Pyruvate dehydrogenase complex, lipoyl-containing component X; E3-binding protein	NM_003477	1.35
collagen, type IV, alpha 1	AI922605	1.35
ankyrin repeat domain 3	NM_020639	1.35
cadherin 3, type 1, P-cadherin (placental)	NM_001793	1.35
butyrophilin, subfamily 1, member A1	NM_001732	1.35
matrix metalloproteinase-like 1	NM_004142	1.35
CUG triplet repeat, RNA binding protein 2	NM_006561	1.35
Ewing sarcoma breakpoint region 1	AL031186	1.35
Consensus includes gb:U88358.1 /DEF=Homo sapiens clone R2 ErbB-3 R2 (c-erbB-3) mRNA, partial cds. /FEA=mRNA /GEN=c-erbB-3 /PROD=ErbB-3 R2 /DB_XREF=gi:3323390 /UG=Hs.3(	U88358	1.35
cytochrome P450, subfamily IIE (ethanol-inducible)	J02843	1.35
histamine H4 receptor	AF312230	1.35
fibroblast growth factor 7 (keratinocyte growth factor)	NM_002009	1.35
paired box gene 5 (B-cell lineage specific activator protein)	NM_016734	1.35
phosphoenolpyruvate carboxytinase 2 (mitochondrial)	NM_004563	1.35
Consensus includes gb:AW979018 /FEA=EST /DB_XREF=gi:8170301 /DB_XREF=est:EST391128 /UG=Hs.132670 ESTs	AW979018	1.35
cytochrome P450, subfamily IVF, polypeptide 8	AF133298	1.35
phosphodiesterase 4D interacting protein (myomegalin)	AI821791	1.35
sperm protein associated with the nucleus, X chromosome, family member A1	NM_013453	1.35
annexin A2 pseudogene 3	M62895	1.35
PBX/knotted 1 homeobox 2	AK023792	1.35
KIAA0530 protein	AA972711	1.35
retinoic acid receptor, gamma	L12060	1.35
Consensus includes gb:AF052145.1 /DEF=Homo sapiens clone 24400 mRNA sequence. /FEA=mRNA /DB_XREF=gi:3360454 /UG=Hs.159528 Homo sapiens clone 24400 mRNA sequence	AF052145	1.35
Consensus includes gb:H94842 /FEA=EST /DB_XREF=gi:1102475 /DB_XREF=est:yu57a03.s1 /CLONE=IMAGE:230188 /UG=Hs.17882 ESTs	H94842	1.35
wingless-type MMTV integration site family, member 11	NM_004626	1.35
retinoic acid receptor, gamma	H57707	1.35
serine (or cysteine) proteinase inhibitor, clade B (ovalbumin), member 13	AI001696	1.35
tenascin XB	NM_019105	1.35
Consensus includes gb:AI275413 /DEF=Homo sapiens partial IGVH3 DP29 gene for immunoglobulin heavy chain V region, case 1, cell Mo VII 116 /FEA=CDS /DB_XREF=gi:7573089 /UG=Hs	AI275413	1.35
natriuretic peptide receptor B/guanylate cyclase B (atrionatriuretic peptide receptor B)	NM_003995	1.35
KIAA0952 protein	NM_014962	1.35
tectorin alpha	NM_005422	1.35
chondroitin sulfate proteoglycan 2 (versican)	BF218922	1.35
gap junction protein, beta 3, 31kD (connexin 31)	AL121988	1.35
myelin oligodendrocyte glycoprotein	U18800	1.35
glycerol kinase	AA292874	1.35
sterol O-acyltransferase 2	AF059203	1.35
mannosidase, alpha, class 2A, member 2	NM_006122	1.35
tryptase beta 1	AF206666	1.35
platelet-derived growth factor beta polypeptide (simian sarcoma viral (v-sis) oncogene homolog)	Z81010	1.35
serine/arginine repetitive matrix 1	NM_005839	1.35
Consensus includes gb:AI740832 /FEA=EST /DB_XREF=gi:5109120 /DB_XREF=est:wg24g02.x1 /CLONE=IMAGE:2366066 /UG=Hs.12311 Homo sapiens clone 23570 mRNA sequence	AI740832	1.35
ATP synthase, H+ transporting, mitochondrial F1 complex, alpha subunit, isoform 1, cardiac muscle	AU140931	1.35

a disintegrin and metalloproteinase domain 17 (tumor necrosis factor, alpha, converting enzyme)	U86755	1.35
glycerol kinase	X68285	1.35
myosin binding protein C, slow type	BF593509	1.35
beta-galactose-3-O-sulfotransferase, 4	NM_024637	1.35
leukemia inhibitory factor (cholinergic differentiation factor)	NM_002309	1.35
integrin, beta 4	NM_000213	1.35
LIM homeobox protein 2	U11701	1.35
c-fos induced growth factor (vascular endothelial growth factor D)	NM_004469	1.35
UDP-glucose ceramide glucosyltransferase-like 2	NM_020121	1.35
Gardner-Rasheed feline sarcoma viral (v-fgr) oncogene homolog	NM_005248	1.35
deiodinase, iodothyronine, type II	AB041843	1.35
apolipoprotein L, 3	NM_014349	1.35
MAGE-like 2	NM_019066	1.35
prostaglandin E receptor 4 (subtype EP4)	AA897516	1.35
carboxypeptidase A4	NM_016352	1.35
hypothetical protein FLJ11726	AL136545	1.35
solute carrier family 17 (sodium phosphate), member 2	NM_005835	1.35
Mdm4, transformed 3T3 cell double minute 4, p53 binding protein (mouse)	NM_002393	1.35
complement component 1, q subcomponent binding protein	M69039	1.35
interleukin 18 receptor accessory protein	NM_003853	1.35
hypothetical protein FLJ12610	NM_024782	1.35
G6B protein	NM_025260	1.35
ribosomal protein S28	BF431363	1.35
zyxtin related factor 1	AI338837	1.35
KIAA1451 protein	BF970829	1.35
down-regulator of transcription 1, TBP-binding (negative cofactor 2)	AW516932	1.35
hypothetical protein FLJ23594	NM_024781	1.35
zinc finger protein 198	AL136621	1.35
cholinergic receptor, nicotinic, alpha polypeptide 4	X89745	1.35
UDP-Gal:betaGlcNAc beta 1,3-galactosyltransferase, polypeptide 4	AB026730	1.35
cytochrome b-245, beta polypeptide (chronic granulomatous disease)	NM_000397	1.35
RCE1 homolog, prenyl protein protease (S. cerevisiae)	AF121951	1.35
zyxin	NM_003461	1.35
serine (or cysteine) proteinase inhibitor, clade B (ovalbumin), member 13	AF169949	1.35
sodium channel, voltage gated, type VIII, alpha polypeptide	NM_014191	1.35
KIAA1036 protein	AV705485	1.35
heparan sulfate (glucosamine) 3-O-sulfotransferase 1	BF000296	1.35
hypothetical protein GL012	NM_030814	1.35
apolipoprotein E	AI358867	1.35
Down syndrome cell adhesion molecule	AF023450	1.35
interleukin 8	AF043337	1.35
hypothetical protein FLJ11560	NM_025182	1.35
endoglin (Osler-Rendu-Weber syndrome 1)	NM_000118	1.35
Rhesus blood group, CcEe antigens	AF312679	1.35
gb:AF130093.1 /DEF=Homo sapiens clone FLB9835 PRO2660 mRNA, complete cds. /FEA=mRNA /PROD=PRO2660 /DB_XREF=gi:11493490 /UG=Hs.302153 Homo sapiens clone FLB9835	AF130093	1.35
CD59 antigen p18-20 (antigen identified by monoclonal antibodies 16.3A5, EJ16, EJ30, EL32 and G344)	NM_000611	1.35
dual specificity phosphatase 3 (vaccinia virus phosphatase VH1-related)	NM_004090	1.35
XAGE-1 protein	NM_020411	1.35
optic atrophy 1 (autosomal dominant)	AB011139	1.35
Consensus includes gb:AK022067.1 /DEF=Homo sapiens cDNA FLJ12005 fis, clone HEMBB1001565. /FEA=mRNA /DB_XREF=gi:10433387 /UG=Hs.287477 Homo sapiens cDNA FLJ12005 f	AK022067	1.35
Consensus includes gb:AK021576.1 /DEF=Homo sapiens cDNA FLJ11514 fis, clone HEMBA1002229. /FEA=mRNA /DB_XREF=gi:10432783 /UG=Hs.306602 Homo sapiens cDNA FLJ11514 f	AK021576	1.35
KIAA1218 protein	AI017382	1.35
hypothetical protein FLJ10748	NM_018203	1.35
Consensus includes gb:BC003535.1 /DEF=Homo sapiens, clone IMAGE:3609326, mRNA, partial cds. /FEA=mRNA /PROD=Unknown (protein for IMAGE:3609326) /DB_XREF=gi:13097635 /L	BC003535	1.35
integrin, alpha 9	NM_002207	1.35
SAR1 protein	BC003658	1.35
ribosomal protein S6 kinase, 90kD, polypeptide 4	NM_003942	1.35
Consensus includes gb:AW969803 /FEA=EST /DB_XREF=gi:8159647 /DB_XREF=est:EST381881 /UG=Hs.105791 ESTs	AW969803	1.35
zinc finger protein (clone 647)	BE350026	1.35
gamma-glutamyltransferase 1	NM_013421	1.35
ankyrin-like with transmembrane domains 1	AA502609	1.35
ATP-binding cassette, sub-family C (CFTR/MRP), member 9	NM_020297	1.35
sphingomyelin phosphodiesterase 1, acid lysosomal (acid sphingomyelinase)	M59916	1.35

KIAA0090 protein	BF197122	1.35
basic transcription element binding protein 1	BF438302	1.35
KIAA0231 protein	R41498	1.35
CD3E antigen, epsilon polypeptide (TIT3 complex)	NM_000733	1.35
solute carrier family 22 (organic anion transporter), member 8	NM_004254	1.35
ubiquitin-conjugating enzyme E2B (RAD6 homolog)	AI768723	1.35
KIAA0495	AI073551	1.35
Consensus includes gb:AV700132 /FEA=EST /DB_XREF=gi:10302103 /DB_XREF=est:AV700132 /CLONE=GKCGSE03 /UG=Hs.295923 seven in absentia (Drosophila) homolog 1	AV700132	1.35
fibrousheathin II	NM_012189	1.35
Consensus includes gb:AA865601 /FEA=EST /DB_XREF=gi:2957877 /DB_XREF=est:og93e03.s1 /CLONE=IMAGE:1455868 /UG=Hs.126261 Homo sapiens Chromosome 16 BAC clone CIT98	AA865601	1.35
guanine nucleotide binding protein (G protein), alpha 11 (Gq class)	NM_002067	1.35
vinculin	AA156675	1.35
corticotropin releasing hormone	NM_000756	1.35
ryanodine receptor 2 (cardiac)	NM_001035	1.35
cytidine deaminase	NM_001785	1.35
phospholipase A2, group VI (cytosolic, calcium-independent)	AF102988	1.35
synovial sarcoma, X breakpoint 2	NM_003147	1.35
itchy homolog E3 ubiquitin protein ligase (mouse)	AI989677	1.35
apoptotic protease activating factor	AF248734	1.35
KIAA0469 gene product	NM_014851	1.35
neuroguanin 2	AF303002	1.35
cadherin 6, type 2, K-cadherin (fetal kidney)	AU151483	1.35
S100 calcium binding protein A2	NM_005978	1.35
GW128 protein	NM_014052	1.35
protein tyrosine phosphatase, non-receptor type substrate 1	AC004832	1.35
platelet factor 4	NM_002619	1.35
v-mos Moloney murine sarcoma viral oncogene homolog	NM_005372	1.35
KIAA1075 protein	AB028998	1.35
hypothetical protein FLJ20623	NM_017905	1.35
Consensus includes gb:U92027.1 /DEF=Human clone 61501 defective mariner transposon Hsmar2 mRNA sequence. /FEA=mRNA /DB_XREF=gi:2052469 /UG=Hs.159211 Human clone 61	U92027	1.35
erythropoietin receptor	M60459	1.35
KIAA0275 gene product	NM_014767	1.35
CDK2-associated protein 1	NM_004642	1.35
sentrin/SUMO-specific protease 3	AK000923	1.35
achaete-scute complex-like 1 (Drosophila)	BC002341	1.35
hypothetical protein MGC20727	BF509069	1.35
C-terminal binding protein 2	BC002486	1.35
guanine nucleotide binding protein (G protein), alpha inhibiting activity polypeptide 3	NM_006496	1.35
hypothetical protein PRO2037	NM_018616	1.35
claudin 9	AI701514	1.35
prostaglandin D2 receptor (DP)	AI460323	1.35
SA (rat hypertension-associated) homolog	NM_005622	1.35
Consensus includes gb:AK025072.1 /DEF=Homo sapiens cDNA: FLJ21419 fis, clone COL04084. /FEA=mRNA /DB_XREF=gi:10437512 /UG=Hs.306759 Homo sapiens cDNA: FLJ21419 fis, cl	AK025072	1.35
DKFZP586L151 protein	AW007573	1.35
KIAA0492 protein	R02172	1.35
Consensus includes gb:AL109678.1 /DEF=Homo sapiens mRNA full length insert cDNA clone EUROIMAGE 28993. /FEA=mRNA /DB_XREF=gi:5689803 /UG=Hs.21597 Homo sapiens mRNA f	AL109678	1.35
development and differentiation enhancing factor 2	NM_003887	1.35
cadherin 6, type 2, K-cadherin (fetal kidney)	NM_004932	1.35
parathyroid hormone receptor 2	NM_005048	1.35
zinc finger protein 84 (HPF2)	NM_003428	1.35
tenascin R (restrictin, janusin)	NM_003285	1.35
cyflicin, basic protein of sperm head cytoskeleton 1	Z22780	1.35
hypothetical protein FLJ12015	NM_024812	1.35
gamma-glutamyltransferase 1	L20492	1.35
protein tyrosine phosphatase, receptor type, f polypeptide (PTPRF), interacting protein (liprin), alpha 2	NM_003625	1.35
a disintegrin-like and metalloprotease (reprolysin type) with thrombospondin type 1 motif, 12	NM_030955	1.35
frizzled homolog 9 (Drosophila)	NM_003508	1.35
Kelch motif containing protein	NM_014458	1.35
major histocompatibility complex, class II, DR beta 3	AA807056	1.35
likely ortholog of mouse NPC derived proline rich protein 1	NM_018212	1.35
adaptor-related protein complex 1, gamma 1 subunit	AL050025	1.35
hypothetical protein FLJ20647	NM_017918	1.35
adrenergic, beta-2-, receptor, surface	NM_000024	1.35

loss of heterozygosity, 11, chromosomal region 2, gene A	NM_014622	1.35
hypothetical protein FLJ20435	NM_017821	1.35
cytochrome b-245, beta polypeptide (chronic granulomatous disease)	AJ308863	1.35
olfactory receptor, family 2, subfamily B, member 2	AJ302584	1.35
interferon regulatory factor 5	AJ028035	1.35
wingless-type MMTV integration site family member 2	NM_003391	1.35
protein tyrosine phosphatase, receptor type, E	AA775177	1.35
hepatitis delta antigen-interacting protein A	AJ583181	1.35
CSH-type zinc finger protein; similar to D. melanogaster muscleblind B protein	AJ088145	1.35
nuclear protein, marker for differentiated aortic smooth muscle and down-regulated with vascular injury	NM_005876	1.35
KIAA1535 protein	BF110802	1.34
immunoglobulin kappa constant	BG540628	1.34
hypothetical protein PRO2730	AJ890972	1.34
MAD, mothers against decapentaplegic homolog 6 (Drosophila)	AJ628464	1.34
chondromodulin I precursor	NM_007015	1.34
small EDRK-rich factor 1B (centromeric)	NM_022978	1.34
runt-related transcription factor 1 (acute myeloid leukemia 1; aml1 oncogene)	L34598	1.34
DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 21	NM_004728	1.34
ectodermal dysplasia 1, anhidrotic	AF061194	1.34
Consensus includes gb:BF970253 /FEA=EST /DB_XREF=gi:12337468 /DB_XREF=est:602273505F1 /CLONE=IMAGE:4361569 /UG=Hs.173684 Homo sapiens mRNA; cDNA DKFZp762G207	BF970253	1.34
KIAA0173 gene product	NM_014640	1.34
cysteine and glycine-rich protein 3 (cardiac LIM protein)	NM_003476	1.34
KIAA0795 protein	NM_025010	1.34
zinc finger protein 23 (KOX 16)	AL567808	1.34
kallikrein B, plasma (Fletcher factor) 1	NM_000892	1.34
hypothetical protein DKFZp434E2220	NM_017612	1.34
member of MYST family histone acetyl transferases, homolog of Drosophila MOF	AL050395	1.34
hypothetical protein FLJ12193	NM_024984	1.34
limbic system-associated membrane protein	NM_002338	1.34
G protein-coupled receptor 19	NM_006143	1.34
KIAA0967 protein	NM_014907	1.34
Consensus includes gb:AF070577.1 /DEF=Homo sapiens clone 24461 mRNA sequence. /FEA=mRNA /DB_XREF=gi:3387949 /UG=Hs.4817 Homo sapiens clone 24461 mRNA sequence	AF070577	1.34
crystallin, beta A4	NM_001886	1.34
hypothetical protein	NM_019062	1.34
guanine nucleotide binding protein (G protein), alpha inhibiting activity polypeptide 1	AL049933	1.34
zinc finger protein 106	NM_022473	1.34
Consensus includes gb:AI884814 /FEA=EST /DB_XREF=gi:5589978 /DB_XREF=est:wi85a05.x1 /CLONE=IMAGE:2431664 /UG=Hs.21964 ESTs	AI884814	1.34
myosin, light polypeptide 5, regulatory	NM_002477	1.34
Consensus includes gb:BF056029 /FEA=EST /DB_XREF=gi:10809925 /DB_XREF=est:7k07f03.x1 /CLONE=IMAGE:3443836 /UG=Hs.174910 ESTs, Weakly similar to dysferlin H.sapiens	BF056029	1.34
hypothetical protein FLJ20551	NM_017875	1.34
nuclear receptor binding factor-2	AA883074	1.34
lymphocyte antigen 117	AF031136	1.34
protocadherin alpha 2	BC003126	1.34
Consensus includes gb:AU145806 /FEA=EST /DB_XREF=gi:11007327 /DB_XREF=est:AU145806 /CLONE=HEMBA1005894 /UG=Hs.296664 Homo sapiens cDNA FLJ11776 fis, clone HEMBA	AU145806	1.34
Consensus includes gb:AF107846 /DEF=Homo sapiens neuroendocrine-specific Golgi protein p55 (XLalphas) gene, exon XL2 and complete cds /FEA=mRNA /DB_XREF=gi:4206779 /UG=Hs	AF107846	1.34
hypothetical protein PRO2949	NM_018544	1.34
leukocyte tyrosine kinase	X52213	1.34
KIAA1416 protein	NM_017780	1.34
cysteinyl leukotriene CysLT2 receptor; cDNA: PSEC0146 from clone PLACE1006979	NM_020377	1.34
iroquois homeobox protein 4	NM_016358	1.34
aquaporin 2 (collecting duct)	NM_000486	1.34
F-box only protein 21	AK001699	1.34
Consensus includes gb:AL117447.1 /DEF=Homo sapiens mRNA; cDNA DKFZp586A0617 (from clone DKFZp586A0617). /FEA=mRNA /DB_XREF=gi:5911895 /UG=Hs.306342 Homo sapiens	AL117447	1.34
Consensus includes gb:AK022063.1 /DEF=Homo sapiens cDNA FLJ12001 fis, clone HEMBB1001535. /FEA=mRNA /DB_XREF=gi:10433383 /UG=Hs.278412 Homo sapiens cDNA FLJ12001 f	AK022063	1.34
secretin	NM_021920	1.34
alpha-1-microglobulin/bikunin precursor	AV645756	1.34
sialidase 3 (membrane sialidase)	NM_006656	1.34
guanine nucleotide binding protein (G protein), beta polypeptide 3	NM_002075	1.34
Fzr1 protein	AJ732960	1.34
Consensus includes gb:AI824954 /FEA=EST /DB_XREF=gi:5445625 /DB_XREF=est:wb04c07.x1 /CLONE=IMAGE:2304684 /UG=Hs.157429 SRY (sex determining region Y)-box 3 /FL=gb:NI	AI824954	1.34
KIAA0582 protein	BG530481	1.34
homeo box C13	NM_017410	1.34
hemoglobin, beta	AF349114	1.34

urotensin 2	NM_021995	1.34
membrane protein, palmitoylated 3 (MAGUK p55 subfamily member 3)	NM_001932	1.34
Consensus includes gb:AK022363.1 /DEF=Homo sapiens cDNA FLJ12301 fis, clone MAMMA1001858. /FEA=mRNA /DB_XREF=gi:10433745 /UG=Hs.287511 Homo sapiens cDNA FLJ12301	AK022363	1.34
mucin 16	NM_024690	1.34
cadherin, EGF LAG seven-pass G-type receptor 2 (flamingo homolog, Drosophila)	D87469	1.34
T cell receptor gamma locus	AC006033	1.34
AF15q14 protein	NM_020380	1.34
T cell receptor gamma constant 2	M13231	1.34
heterogeneous nuclear ribonucleoprotein U (scaffold attachment factor A)	D13413	1.34
peptidylprolyl isomerase E (cyclophilin E)	BF511556	1.34
hypothetical protein FLJ20509	NM_017851	1.34
hypothetical protein FLJ10339	NM_018063	1.34
TGFB-induced factor 2 (TALE family homeobox)	NM_021809	1.34
mucin 3A, intestinal	AF007194	1.34
chromosome 21 open reading frame 68	NM_024944	1.34
adaptor-related protein complex 4, sigma 1 subunit	BC001259	1.34
myosin IA	AF009961	1.34
carboxyl ester lipase-like (bile salt-stimulated lipase-like)	NM_001808	1.34
Consensus includes gb:AA716107 /FEA=EST /DB_XREF=gi:2728381 /DB_XREF=est:zg63d06.s1 /CLONE=IMAGE:398027 /UG=Hs.8963 Homo sapiens mRNA full length insert cDNA clone EI	AA716107	1.34
MYA22 protein	BF590317	1.34
human immunodeficiency virus type I enhancer binding protein 1	NM_002114	1.34
hepatocyte nuclear factor 4, alpha	NM_000457	1.34
Consensus includes gb:AL049233.1 /DEF=Homo sapiens mRNA; cDNA DKFZp564A023 (from clone DKFZp564A023). /FEA=mRNA /DB_XREF=gi:4499967 /UG=Hs.42244 Homo sapiens mRI	AL049233	1.34
matrix metalloproteinase 14 (membrane-inserted)	Z48481	1.34
forkhead box F2	NM_001452	1.34
son of sevenless homolog 1 (Drosophila)	AA700167	1.34
interleukin 11	M57765	1.34
LIM domain kinase 1	NM_002314	1.34
transcription factor 3 (E2A immunoglobulin enhancer binding factors E12/E47)	AI871234	1.34
hypothetical protein FLJ20244	AL390133	1.34
transcription elongation factor B (SIII), polypeptide 1-like	BE964043	1.34
zinc finger protein	AA349848	1.34
gonadotropin-regulated testicular RNA helicase	NM_013264	1.34
hypothetical protein FLJ23282	NM_024816	1.34
signal transducer and activator of transcription 1, 91kD	BC002704	1.34
keratin, hair, acidic, 8	NM_006771	1.34
Consensus includes gb:AA156865 /FEA=EST /DB_XREF=gi:1728480 /DB_XREF=est:zi20f08.s1 /CLONE=IMAGE:502503 /UG=Hs.22370 Homo sapiens mRNA; cDNA DKFZp564D0122 (from	AA156865	1.34
hypothetical protein FLJ10634	NM_018163	1.34
Consensus includes gb:AK023895.1 /DEF=Homo sapiens cDNA FLJ13833 fis, clone THYRO1000676. /FEA=mRNA /DB_XREF=gi:10435969 /UG=Hs.296745 Homo sapiens cDNA FLJ13833 fi	AK023895	1.34
Consensus includes gb:AI439416 /FEA=EST /DB_XREF=gi:4304169 /DB_XREF=est:tj04b04.x1 /CLONE=IMAGE:2140495 /UG=Hs.165728 ESTs	AI439416	1.34
calcium channel, voltage-dependent, alpha 1G subunit	AB012043	1.34
transcription factor AP-2 alpha (activating enhancer binding protein 2 alpha)	BF343007	1.34
G protein-coupled receptor, family C, group 5, member B	NM_016235	1.34
KIAA0246 protein	D87433	1.34
Consensus includes gb:Y10259.1 /DEF=H.sapiens ACTH receptor mRNA 3UTR. /FEA=mRNA /DB_XREF=gi:2230860 /UG=Hs.274501 H.sapiens ACTH receptor mRNA 3UTR	Y10259	1.34
hypothetical protein PRO0889	NM_018496	1.34
receptor (calcitonin) activity modifying protein 1	NM_005855	1.34
golgi SNAP receptor complex member 2	AW149492	1.34
hypothetical protein PP1665	AL041124	1.34
casein, kappa	NM_005212	1.34
activin A receptor, type IB	NM_020328	1.34
mutated in colorectal cancers	NM_002387	1.34
hypothetical protein FLJ11715	NM_024564	1.34
sonic hedgehog homolog (Drosophila)	NM_000193	1.34
Consensus includes gb:AI057637 /FEA=EST /DB_XREF=gi:3331503 /DB_XREF=est:oy31h06.x1 /CLONE=IMAGE:1667483 /UG=Hs.234898 ESTs, Weakly similar to 2109260A B cell growth I	AI057637	1.34
CGI-148 protein	NM_016078	1.34
Consensus includes gb:AW973235 /FEA=EST /DB_XREF=gi:8163081 /DB_XREF=est:EST385333 /UG=Hs.293697 ESTs	AW973235	1.34
kallikrein 11	NM_006853	1.34
homogentisate 1,2-dioxygenase (homogentisate oxidase)	AI478172	1.34
matrix Gla protein	NM_000900	1.34
Consensus includes gb:U80773.1 /DEF=Human EST clone 42944 mariner transposon Hsmar1 sequence. /FEA=mRNA /DB_XREF=gi:2231376 /UG=Hs.121580 Human EST clone 42944 mari	U80773	1.34
Consensus includes gb:AK025363.1 /DEF=Homo sapiens cDNA: FLJ21710 fis, clone COL10087. /FEA=mRNA /DB_XREF=gi:10437865 /UG=Hs.306807 Homo sapiens cDNA: FLJ21710 fis, cl	AK025363	1.34
putative homeodomain transcription factor 1	AA927671	1.34

Meis1, myeloid ecotropic viral integration site 1 homolog (mouse)	NM_002398	1.34
pre-T/NK cell associated protein	NM_004908	1.34
complement component 1, q subcomponent, receptor 1	W72082	1.34
eukaryotic translation initiation factor 3, subunit 1 (alpha, 35kD)	NM_003758	1.34
phospholipase A2 receptor 1, 180kD	NM_007366	1.34
growth arrest-specific 7	BC006454	1.34
troponin I, skeletal, fast	NM_003282	1.34
KIAA0534 protein	AW151108	1.34
leucine zipper, putative tumor suppressor 1	N21184	1.34
hypothetical protein FLJ12168	NM_024682	1.34
likely ortholog of mouse zinc finger homeodomain 4	NM_024721	1.34
dihydrolipoamide branched chain transacylase (E2 component of branched chain keto acid dehydrogenase complex; maple syrup urine disease)	M19301	1.34
RAB6A, member RAS oncogene family	BC003617	1.34
collagen, type V, alpha 3	NM_015719	1.34
solute carrier family 25 (mitochondrial oxodicarboxylate carrier), member 21	NM_030631	1.34
laminin, gamma 3	NM_006059	1.34
uncharacterized hypothalamus protein HT010	AV716798	1.34
myelin protein zero (Charcot-Marie-Tooth neuropathy 1B)	D10537	1.34
phosphoglycerate mutase 2 (muscle)	NM_000290	1.34
hypothetical protein FLJ10921	NM_018272	1.34
hypothetical protein	NM_016626	1.34
CGI-35 protein	NM_015962	1.34
collagen, type XIII, alpha 1	NM_005203	1.34
KIAA1085 protein	AU160676	1.34
insulin-like growth factor binding protein 5	AW157548	1.34
B-cell CLL/lymphoma 10	AF082283	1.34
Consensus includes gb:AF052176.1 /DEF=Homo sapiens clone 24457 mRNA sequence. /FEA=mRNA /DB_XREF=gi:3360487 /UG=Hs.158529 Homo sapiens clone 24457 mRNA sequence	AF052176	1.34
KIAA0551 protein	R59093	1.34
Consensus includes gb:J04178.1 /DEF=Human abnormal beta-hexosaminidase alpha chain (HEXA) mRNA, partial cds. /FEA=mRNA /DB_XREF=gi:184001 /UG=Hs.166299 Human abnormal	J04178	1.34
N-acetylglucosamine-phosphate mutase	AV727934	1.34
MAD, mothers against decapentaplegic homolog 4 (Drosophila)	U44378	1.34
desmocollin 2	NM_004949	1.34
runt-related transcription factor 1 (acute myeloid leukemia 1; aml1 oncogene)	NM_001754	1.34
cold autoinflammatory syndrome 1	NM_004895	1.34
cholinergic receptor, muscarinic 2	NM_000739	1.34
helicase-moi	AK001827	1.34
Consensus includes gb:AW008921 /FEA=EST /DB_XREF=gi:5857699 /DB_XREF=est:ws75a03.x1 /CLONE=IMAGE:2503756 /UG=Hs.13138 Homo sapiens, clone IMAGE:3448343, mRNA, pa	AW008921	1.34
troponin C2, fast	NM_003279	1.34
A kinase (PRKA) anchor protein 7	NM_004842	1.34
hypothetical protein FLJ20739	NM_017949	1.34
proline-rich protein BstNI subfamily 2	K03205	1.34
mannan-binding lectin serine protease 1 (C4/C2 activating component of Ra-reactive factor)	NM_001879	1.34
TRF-proximal protein	NM_004275	1.34
Consensus includes gb:AL049443.1 /DEF=Homo sapiens mRNA; cDNA DKFZp586N2020 (from clone DKFZp586N2020). /FEA=mRNA /DB_XREF=gi:4500224 /UG=Hs.161283 Homo sapiens	AL049443	1.34
zinc finger protein 148 (pHZ-52)	NM_021964	1.34
eve, even-skipped homeo box homolog 1 (Drosophila)	NM_001989	1.34
Consensus includes gb:AU147317 /FEA=EST /DB_XREF=gi:11008838 /DB_XREF=est:AU147317 /CLONE=MAMMA1000340 /UG=Hs.287491 Homo sapiens cDNA FLJ12140 fis, clone MAMM	AU147317	1.34
tissue inhibitor of metalloproteinase 3 (Sorsby fundus dystrophy, pseudoinflammatory)	AW338933	1.34
hypothetical protein FLJ10094	NM_017993	1.34
Consensus includes gb:AL049544 /DEF=Human DNA sequence from clone 24088 on chromosome 6p11.2-q12. Contains the 3 part of a gene for a novel protein similar to T-STAR, Etoile, Sa	AL049544	1.34
v-ski sarcoma viral oncogene homolog (avian)	BF431501	1.34
hypothetical protein MGC4293	AI370867	1.34
DNA-binding protein amplifying expression of surfactant protein B	NM_006692	1.34
tumor necrosis factor (ligand) superfamily, member 11	AB037599	1.34
lymphocyte-specific protein 1	NM_002339	1.34
SH3-domain GRB2-like 3	NM_003027	1.34
hypothetical protein FLJ11294	NM_018383	1.34
serum/glucocorticoid regulated kinase	NM_005627	1.34
protein phosphatase 1, regulatory subunit 10	AI492873	1.34
hypothetical protein FLJ20132	NM_017682	1.34
phosphoinositide-3-kinase, regulatory subunit, polypeptide 3 (p55, gamma)	AF028785	1.34
DKFZP586M2123 protein	AI671186	1.34
hypothetical protein	AA835004	1.34

inducible T-cell co-stimulator	AB023135	1.34
KIAA1128 protein	AF241785	1.34
glutamate receptor, metabotropic 7	X94552	1.34
synovial sarcoma, X breakpoint 2	AF257500	1.34
glutaminase	AF097495	1.34
hypothetical protein MGC4170	NM_024312	1.34
hypothetical protein from clone 643	BC003379	1.34
hypothetical protein FLJ25965	AC002398	1.34
stratifin	X57348	1.34
pregnancy specific beta-1-glycoprotein 1	NM_006905	1.34
similar to rat integral membrane glycoprotein POM121	AU158490	1.34
hypothetical protein FLJ11316	NM_018388	1.34
rearranged L-myc fusion sequence	NM_012421	1.34
chromosome 21 open reading frame 2	AI435747	1.34
sprouty (Drosophila) homolog 4	NM_030964	1.34
synuclein, gamma (breast cancer-specific protein 1)	AF010126	1.34
syntrophin, gamma 1	NM_018967	1.34
potassium voltage-gated channel, shaker-related subfamily, member 3	NM_002232	1.34
aspartate beta-hydroxylase	NM_004318	1.34
KIAA0469 gene product	AL031447	1.34
protocadherin gamma subfamily A, 8	AB002325	1.34
corticotropin releasing hormone receptor 2	AF019381	1.34
dihydropyrimidine dehydrogenase	NM_000110	1.34
cer-d4 (mouse) homolog	NM_012074	1.34
dishevelled, dsh homolog 2 (Drosophila)	AW016304	1.34
cytochrome P450, subfamily IIIA (niphedipine oxidase), polypeptide 5	NM_000777	1.34
promyelocytic leukemia	M79462	1.34
Consensus includes gb:AL096700 /DEF=Human DNA sequence from clone RP3-499B10 on chromosome Xp22.31-22.13. Contains the 3 end of the gene for serinethreonine protein phospha	AL096700	1.34
interleukin 17 (cytotoxic T-lymphocyte-associated serine esterase 8)	NM_002190	1.34
Consensus includes gb:AK001619.1 /DEF=Homo sapiens cDNA FLJ10757 fis, clone NT2RP3004578, highly similar to Homo sapiens mRNA for KIAA0477 protein. /FEA=mRNA /DB_XREF=gi:	AK001619	1.34
guanine nucleotide binding protein 11	NM_004126	1.34
immunoglobulin superfamily containing leucine-rich repeat	NM_005545	1.34
bone morphogenetic protein 4	D30751	1.34
Consensus includes gb:AI983904 /FEA=EST /DB_XREF=gi:5811123 /DB_XREF=est:wz55d05.x1 /CLONE=IMAGE:2561961 /UG=Hs.79732 fibulin 1	AI983904	1.34
Consensus includes gb:AK022065.1 /DEF=Homo sapiens cDNA FLJ12003 fis, clone HEMBB1001537. /FEA=mRNA /DB_XREF=gi:10433385 /UG=Hs.296683 Homo sapiens cDNA FLJ12003 f	AK022065	1.34
solute carrier family 19 (folate transporter), member 1	AF004354	1.34
cytochrome P450, subfamily IIIA (niphedipine oxidase), polypeptide 4	AF182273	1.34
KIAA0101 gene product	BC005832	1.34
Consensus includes gb:AK027254.1 /DEF=Homo sapiens cDNA: FLJ23601 fis, clone LNG15501. /FEA=mRNA /DB_XREF=gi:10440343 /UG=Hs.306918 Homo sapiens cDNA: FLJ23601 fis, cl	AK027254	1.34
peroxisome proliferative activated receptor, gamma	NM_015869	1.34
interferon, alpha 5	NM_002169	1.34
integrin, beta 3 (platelet glycoprotein IIIa, antigen CD61)	AI151479	1.34
Consensus includes gb:AL137713.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434I0523 (from clone DKFZp434I0523). /FEA=mRNA /DB_XREF=gi:6808127 /UG=Hs.306465 Homo sapiens n	AL137713	1.34
hypothetical protein FLJ10142	NM_018008	1.34
hypothetical protein FLJ23239	NM_022766	1.34
KIAA1117 protein	AB029040	1.34
5-hydroxytryptamine (serotonin) receptor 1E	NM_000865	1.34
oxidative 3 alpha hydroxysteroid dehydrogenase; retinol dehydrogenase; 3-hydroxysteroid epimerase	NM_003725	1.34
Consensus includes gb:BF062299 /FEA=EST /DB_XREF=gi:10821197 /DB_XREF=est:7k76c03.x1 /CLONE=IMAGE:3481325 /UG=Hs.323079 Homo sapiens mRNA; cDNA DKFZp564P116 (fr	BF062299	1.34
superoxide dismutase 3, extracellular	NM_003102	1.34
N-myc downstream regulated	NM_006096	1.34
HycD family inhibitor	NM_005586	1.34
65K protein	NM_025262	1.34
postmeiotic segregation increased 2-like 8	AB017005	1.33
natriuretic peptide receptor A/guanylate cyclase A (atrionatriuretic peptide receptor A)	X15357	1.33
fibrinogen-like 1	NM_004467	1.33
isocitrate dehydrogenase 2 (NADP+), mitochondrial	AU151428	1.33
phospholipase A2, group VII (platelet-activating factor acetylhydrolase, plasma)	NM_005084	1.33
deleted in lung and esophageal cancer 1	NM_007337	1.33
CUG triplet repeat, RNA binding protein 2	U69546	1.33
DKFZP434N093 protein	AL080144	1.33
HIC1-related gene on chromosome 22	AL162003	1.33
collagen, type V, alpha 2	NM_000393	1.33

connector enhancer of KSR-like (Drosophila kinase suppressor of ras)	NM_006314	1.33
solute carrier family 16 (monocarboxylic acid transporters), member 3	AA527578	1.33
cell division cycle 25C	AF277724	1.33
hypothetical protein FLJ11068	NM_018314	1.33
junctional adhesion molecule 2	NM_021219	1.33
dishevelled, dsh homolog 2 (Drosophila)	NM_004422	1.33
Consensus includes gb:BE669609 /FEA=EST /DB_XREF=gi:10030150 /DB_XREF=est:7e15h11.x1 /CLONE=IMAGE:3282597 /UG=Hs.251411 Homo sapiens mRNA; cDNA DKFZp566B233 (h	BE669609	1.33
Consensus includes gb:AL359563.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434D0326 (from clone DKFZp434D0326). /FEA=mRNA /DB_XREF=gi:8655619 /UG=Hs.306509 Homo sapiens	AL359563	1.33
adenylate cyclase 9	AB011092	1.33
CUG triplet repeat, RNA binding protein 2	N36839	1.33
cytochrome P450, subfamily IIA (phenobarbital-inducible), polypeptide 6	AF182275	1.33
likely ortholog of mouse testis expressed gene 27	NM_021943	1.33
biglycan	NM_001711	1.33
small inducible cytokine A4	NM_002984	1.33
transcriptional unit N143	AI002572	1.33
hypothetical protein FLJ20363	NM_017784	1.33
CD24 antigen (small cell lung carcinoma cluster 4 antigen)	BG327863	1.33
axonal transport of synaptic vesicles	NM_004321	1.33
thyroid hormone receptor-associated protein, 240 kDa subunit	AB011165	1.33
dual-specificity tyrosine-(Y)-phosphorylation regulated kinase 2	AI192838	1.33
calpastatin	BG111635	1.33
armadillo repeat gene deletes in velocardiofacial syndrome	NM_001670	1.33
myelin associated glycoprotein	X98405	1.33
Consensus includes gb:AF070628.1 /DEF=Homo sapiens clone 24803 mRNA sequence. /FEA=mRNA /DB_XREF=gi:3283895 /UG=Hs.32766 Homo sapiens clone 24803 mRNA sequence	AF070628	1.33
Consensus includes gb:AF043584.1 /DEF=Homo sapiens clone ASMneg1-b1 immunoglobulin lambda chain VI region, (IGL) mRNA, partial cds. /FEA=mRNA /GEN=IGL /PROD=immunoglobul	AF043584	1.33
Consensus includes gb:AA131324 /FEA=EST /DB_XREF=gi:1693011 /DB_XREF=est:zl31b04.s1 /CLONE=IMAGE:503503 /UG=Hs.28264 Homo sapiens mRNA; cDNA DKFZp564L0822 (from	AA131324	1.33
gb:AF116705.1 /DEF=Homo sapiens PRO2489 mRNA, complete cds. /FEA=mRNA /PROD=PRO2489 /DB_XREF=gi:7959908 /FL=gb:AF116705.1	AF116705	1.33
Consensus includes gb:AV712602 /FEA=EST /DB_XREF=gi:10731908 /DB_XREF=est:AV712602 /CLONE=DCAACD04 /UG=Hs.5957 Homo sapiens clone 24416 mRNA sequence	AV712602	1.33
Consensus includes gb:AL137270.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434G0614 (from clone DKFZp434G0614). /FEA=mRNA /DB_XREF=gi:6807705 /UG=Hs.39337 Homo sapiens n	AL137270	1.33
ubiquitin associated protein	AL039447	1.33
Consensus includes gb:AB041269.1 /DEF=Homo sapiens mRNA for keratin 19, partial cds, isolate:K19-141. /FEA=mRNA /GEN=keratin 19 /PROD=keratin 19 /DB_XREF=gi:7594735 /UG=H	AB041269	1.33
forkhead box L2	NM_023067	1.33
protein tyrosine phosphatase, non-receptor type 21	X79510	1.33
secretory pathway component Sec31B-1	AF274863	1.33
cadherin, EGF LAG seven-pass G-type receptor 2 (flamingo homolog, Drosophila)	NM_001408	1.33
hypothetical protein FLJ22127	AA203219	1.33
host cell factor 2	NM_013320	1.33
5-hydroxytryptamine (serotonin) receptor 6	AF007141	1.33
glutamate receptor, metabotropic 1	U31216	1.33
indian hedgehog homolog (Drosophila)	BE869172	1.33
Consensus includes gb:AL050335 /DEF=Human DNA sequence from clone RP1-190J20 on chromosome 6p22.3-24.1. Contains ESTs, STSs and GSSs. Contains a putative novel gene and an R	AL050335	1.33
Consensus includes gb:AK025194.1 /DEF=Homo sapiens cDNA: FLJ21541 fis, clone COL06166. /FEA=mRNA /DB_XREF=gi:10437657 /UG=Hs.306784 Homo sapiens cDNA: FLJ21541 fis, cl	AK025194	1.33
Consensus includes gb:AI927067 /FEA=EST /DB_XREF=gi:5663031 /DB_XREF=est:wo87f01.x1 /CLONE=IMAGE:2462329 /UG=Hs.289068 Homo sapiens cDNA FLJ11918 fis, clone HEMBB1	AI927067	1.33
glutamate receptor, metabotropic 6	NM_000843	1.33
microtubule-associated protein like echinoderm EMAP	AF103939	1.33
Consensus includes gb:AW969655 /FEA=EST /DB_XREF=gi:8159499 /DB_XREF=est:EST381732 /UG=Hs.168386 ESTs	AW969655	1.33
aldolase B, fructose-bisphosphate	NM_000035	1.33
gb:NM_020484.1 /DEF=Homo sapiens RAGE binding protein (AF011757), mRNA. /FEA=mRNA /GEN=AF011757 /PROD=RAGE binding protein /DB_XREF=gi:10048411 /UG=Hs.105937 RA	NM_020484	1.33
matrix metalloproteinase 16 (membrane-inserted)	NM_022564	1.33
Consensus includes gb:Y10259.1 /DEF=H.sapiens ACTH receptor mRNA 3UTR. /FEA=mRNA /DB_XREF=gi:2230860 /UG=Hs.274501 H.sapiens ACTH receptor mRNA 3UTR	Y10259	1.33
extracellular matrix protein 2, female organ and adipocyte specific	NM_001393	1.33
KIAA0810 protein	NM_025154	1.33
early growth response 1	NM_001964	1.33
Consensus includes gb:U79248.1 /DEF=Human clone 23826 mRNA sequence. /FEA=mRNA /DB_XREF=gi:1710195 /UG=Hs.12484 Human clone 23826 mRNA sequence	U79248	1.33
sialic acid binding Ig-like lectin 8	NM_014442	1.33
bone morphogenetic protein 10	NM_014482	1.33
NTF2-related export protein 1	NM_013248	1.33
synovial sarcoma, X breakpoint 4	BC005325	1.33
pleckstrin homology, Sec7 and coiled/coil domains 4	NM_013385	1.33
protein O-fucosyltransferase 1	AL045513	1.33
perilipin	NM_002666	1.33
KIAA0551 protein	N25621	1.33
glycoprotein 2 (zymogen granule membrane)	NM_016295	1.33

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KIAA0663 gene product	AI803216	1.33
sarcospan (Kras oncogene-associated gene)	NM_005086	1.33
cathepsin 6	NM_001911	1.33
erythropoietin receptor	X97671	1.33
transcription factor-like 4	AF213666	1.33
putative zinc finger protein NY-REN-34 antigen	BF055474	1.33
adenosine deaminase, RNA-specific, B1 (homolog of rat RED1)	NM_001112	1.33
claudin 14	AF314090	1.33
hypothetical protein FLJ10628	NM_018159	1.33
Consensus includes gb:AI923984 /FEA=EST /DB_XREF=gi:5659948 /DB_XREF=est:wn49d12.x1 /CLONE=IMAGE:2448791 /UG=Hs.46320 Small proline-rich protein SPRK human, odontoger	AI923984	1.33
matrix metalloproteinase 14 (membrane-inserted)	NM_004995	1.33
paired box gene 8	NM_013951	1.33
RAR-related orphan receptor A	L14611	1.33
catechol-O-methyltransferase	BG054676	1.33
hypothetical protein FLJ12681	AI003763	1.33
adenylate cyclase 6	AF250226	1.33
tyrosine aminotransferase	AV647713	1.33
neuronal pentraxin I	NM_002522	1.33
fibrinogen, A alpha polypeptide	NM_021871	1.33
ets variant gene 1	AC004857	1.33
CASP8 and FADD-like apoptosis regulator	AF005775	1.33
hypothetical protein DKFZp434E1723	AI199589	1.33
zinc finger protein 278	AF242522	1.33
heart and neural crest derivatives expressed 2	NM_021973	1.33
prophet of Pit1, paired-like homeodomain transcription factor	AF076215	1.33
nucleosome assembly protein 1-like 3	NM_004538	1.33
protein homologous to salivary proline-rich protein P-B	NM_012390	1.33
pleiomorphic adenoma gene-like 1	BG547855	1.33
epidermal growth factor receptor (erythroblastic leukemia viral (v-erb-b) oncogene homolog, avian)	NM_005228	1.33
CGI-130 protein	NM_016063	1.33
potassium voltage-gated channel, subfamily F, member 1	AF029780	1.33
hypothetical protein ASH1	NM_018489	1.33
Rho GTPase activating protein 8	AA533284	1.33
hypothetical protein FLJ20071	NM_017653	1.33
KIAA1886 protein	AL136572	1.33
chromosome 8 open reading frame 1	NM_004337	1.33
KIAA0557 protein	AI871396	1.33
hypothetical protein FLJ11795	NM_024669	1.33
p21(CDKN1A)-activated kinase 6	AI236915	1.33
SET binding factor 1	AI333707	1.33
chromosome 21 open reading frame 7	NM_020152	1.33
zinc finger protein 174	BC001161	1.33
hypothetical protein FLB3342	NM_020118	1.33
acyl-Coenzyme A dehydrogenase, long chain	NM_001608	1.33
sema domain, immunoglobulin domain (Ig), transmembrane domain (TM) and short cytoplasmic domain, (semaphorin) 4F	AF119878	1.33
neurogenic differentiation 2	AB021742	1.33
dual-specificity tyrosine-(Y)-phosphorylation regulated kinase 1B	NM_004714	1.33
G protein-coupled receptor 64	NM_005756	1.33
CCAAT/enhancer binding protein (C/EBP), delta	NM_005195	1.33
promyelocytic leukemia	BC000080	1.33
serine protease inhibitor, Kazal type 4	NM_014471	1.33
armadillo repeat gene deletes in velocardiofacial syndrome	BG149428	1.33
hypothetical protein FLJ13110	NM_022912	1.33
pleckstrin and Sec7 domain protein	NM_002779	1.33
glycerol kinase pseudogene 2	AA757089	1.33
hypothetical protein FLJ20359	NM_017781	1.33
serine (or cysteine) proteinase inhibitor, clade F (alpha-2 antiplasmin, pigment epithelium derived factor), member 2	NM_000934	1.33
hypothetical gene CG030	U50531	1.33
hypothetical protein FLJ11539	NM_024748	1.33
Consensus includes gb:AK024909.1 /DEF=Homo sapiens cDNA: FLJ21256 fis, clone COL01402. /FEA=mRNA /DB_XREF=gi:10437324 /UG=Hs.306724 Homo sapiens cDNA: FLJ21256 fis, cl	AK024909	1.33
matrix, extracellular phosphoglycoprotein with ASARM motif (bone)	NM_020203	1.33
solute carrier family 4, anion exchanger, member 3	NM_005070	1.33
Homer, neuronal immediate early gene, 3	AI871287	1.33

dystroglycan 1 (dystrophin-associated glycoprotein 1)	AW411370	1.33
dihydropyrimidinase-like 4	NM_006426	1.33
tripartite motif-containing 29	AF230389	1.33
CGI-83 protein	NM_016027	1.33
hOAT4	NM_018484	1.33
phospholipase A2, group IIE	NM_014589	1.33
carnitine acetyltransferase	NM_000755	1.33
potassium voltage-gated channel, KQT-like subfamily, member 2	D82346	1.33
Consensus includes gb:AL049337.1 /DEF=Homo sapiens mRNA; cDNA DKFZp564P016 (from clone DKFZp564P016). /FEA=mRNA /DB_XREF=gi:4500118 /UG=Hs.132571 Homo sapiens ml	AL049337	1.33
Consensus includes gb:AK025044.1 /DEF=Homo sapiens cDNA: FLJ21391 fis, clone COL03479. /FEA=mRNA /DB_XREF=gi:10437481 /UG=Hs.306752 Homo sapiens cDNA: FLJ21391 fis, cl	AK025044	1.33
rabA binding protein 1	AW025261	1.33
mitogen-activated protein kinase kinase kinase 6	NM_004672	1.33
sorting nexin 16	NM_022133	1.33
MAX binding protein	NM_020310	1.33
Consensus includes gb:BF939689 /FEA=EST /DB_XREF=gi:12357100 /DB_XREF=est:nac80b04.x1 /CLONE=IMAGE:3440575 /UG=Hs.159629 myosin IXB	BF939689	1.33
hypothetical protein MGC3048	NM_024052	1.33
GKD01 protein	NM_020198	1.33
Lutheran blood group (Auberger b antigen included)	NM_005581	1.33
serine/threonine kinase 23	NM_014370	1.33
angiopoietin 1	AF209975	1.33
slit homolog 2 (Drosophila)	AF055585	1.33
cAMP responsive element modulator	S68134	1.33
hypothetical protein FLJ12748	BC006434	1.33
NEU1 protein	NM_016215	1.33
thyroid hormone receptor coactivating protein	AL136823	1.33
neurofibromin 1 (neurofibromatosis, von Recklinghausen disease, Watson disease)	D42072	1.33
ubiquitin specific protease 22	AL162082	1.33
COPZ2 for nonclathrin coat protein zeta-COP	NM_016429	1.33
breast cell glutaminase	NM_013267	1.33
myomesin 1 (skelemin) (185kD)	NM_003803	1.33
mutS homolog 4 (E. coli)	AF104243	1.33
proline-rich Gla (G-carboxyglutamic acid) polypeptide 2	NM_000951	1.33
natural killer cell receptor 284	NM_016382	1.33
Consensus includes gb:AL390738 /DEF=Human DNA sequence from clone RP11-438F9 on chromosome 13 Contains the gene for the ortholog of mouse homeobox protein GSH1, the 3 end	AL390738	1.33
glucosidase, alpha; acid (Pompe disease, glycogen storage disease type II)	NM_000152	1.33
zinc finger protein RINZF	NM_023929	1.33
Kallmann syndrome 1 sequence	NM_000216	1.33
proline-rich protein BstNI subfamily 1	NM_005039	1.33
solute carrier family 6 (neurotransmitter transporter, noradrenalin), member 2	BC000563	1.33
FXYD domain-containing ion transport regulator 7	NM_022006	1.33
Consensus includes gb:H15535 /FEA=EST /DB_XREF=gi:880355 /DB_XREF=est:ym27c01.s1 /CLONE=IMAGE:49385 /UG=Hs.52792 Homo sapiens mRNA; cDNA DKFZp586I1823 (from clon	H15535	1.33
cytochrome P450, subfamily IID (debrisoquine, sparteine, etc., -metabolizing), polypeptide 6	X16866	1.33
carcinoembryonic antigen-related cell adhesion molecule 6 (non-specific cross reacting antigen)	BC005008	1.33
likely homolog of mouse immunity-associated nucleotide 4	AI435089	1.33
secreted phosphoprotein 2, 24kD	NM_006944	1.33
ets variant gene 6 (TEL oncogene)	NM_001987	1.33
RAB13, member RAS oncogene family	NM_002870	1.33
zinc finger protein 278	AF254087	1.33
Consensus includes gb:AL136967 /DEF=Human DNA sequence from clone RP1-149M18 on chromosome 6 Contains ESTs, STSs, GSSs and CpG islands. Contains the natural killer cell p44 rela	AL136967	1.33
gap junction protein, alpha 4, 37kD (connexin 37)	M96789	1.33
eukaryotic translation initiation factor 1A, Y chromosome	NM_004681	1.33
cyclin-dependent kinase 5, regulatory subunit 2 (p39)	R51311	1.33
UDP-Gal:betaGlcNAc beta 1,4- galactosyltransferase, polypeptide 1	U10473	1.33
cholinergic receptor, nicotinic, alpha polypeptide 5	NM_000745	1.33
regulator of G-protein signalling 6	AF073921	1.33
Consensus includes gb:AA773554 /FEA=EST /DB_XREF=gi:2825125 /DB_XREF=est:ab61a05.s1 /CLONE=IMAGE:845264 /UG=Hs.234216 Human 40871 mRNA partial sequence	AA773554	1.33
cholinergic receptor, nicotinic, alpha polypeptide 4	L35901	1.33
neuro-oncological ventral antigen 2	AI040329	1.33
UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase 4 (GalNAc-T4)	NM_003774	1.33
dystrophin related protein 2	NM_001939	1.33
chromosome 14 open reading frame 1	NM_007176	1.33
zona pellucida glycoprotein 2 (sperm receptor)	NM_003460	1.33
Consensus includes gb:AV700338 /FEA=EST /DB_XREF=gi:10302309 /DB_XREF=est:AV700338 /CLONE=GKCAMH12 /UG=Hs.286243 ESTs	AV700338	1.33

chromosome 11 open reading frame 11	AB014559	1.33
TIA1 cytotoxic granule-associated RNA binding protein	AL567227	1.33
podocalyxin-like	NM_005397	1.33
EphB2	AL530874	1.33
receptor tyrosine kinase-like orphan receptor 1	NM_005012	1.33
SHT3 suppressor of mif two 3 homolog 2 (yeast)	AJ971724	1.33
protocadherin 16	BF222893	1.33
small nuclear RNA activating complex, polypeptide 4, 190kD	AK023513	1.33
Consensus includes gb:AW516297 /FEA=EST /DB_XREF=gi:7154379 /DB_XREF=est:xt64d08.x1 /CLONE=IMAGE:2791215 /UG=Hs.323231 Homo sapiens cDNA FLJ11946 fis, clone HEMBB	AW516297	1.33
phosphatidylinositol glycan, class C, pseudogene 1	AW305097	1.33
cancer/testis antigen 2	AJ012833	1.33
Consensus includes gb:AL353943.1 /DEF=Homo sapiens mRNA; cDNA DKFZp761E0611 (from clone DKFZp761E0611). /FEA=mRNA /GEN=DKFZp761E0611 /PROD=hypothetical protein /DI	AL353943	1.33
cytochrome P450, subfamily IVF, polypeptide 3 (leukotriene B4 omega hydroxylase)	NM_000896	1.33
KIAA0998 protein	NM_015072	1.33
leucine-zipper-like transcriptional regulator, 1	NM_006767	1.33
cytokine-like protein C17	NM_018659	1.33
parathyroid hormone receptor 1	NM_000316	1.33
zinc finger protein 20 (KOX 13)	AU154474	1.33
carcinoembryonic antigen-related cell adhesion molecule 4	NM_001817	1.33
KIAA0624 protein	AB014524	1.33
hypothetical protein FLJ11457	NM_024753	1.33
a disintegrin-like and metalloprotease (reprolysin type) with thrombospondin type 1 motif, 6	NM_014273	1.33
emopamil binding protein (sterol isomerase)	NM_006579	1.33
Consensus includes gb:AL080107.1 /DEF=Homo sapiens mRNA; cDNA DKFZp586D0522 (from clone DKFZp586D0522). /FEA=mRNA /DB_XREF=gi:5262532 /UG=Hs.163173 Homo sapiens	AL080107	1.33
EphB4	NM_004444	1.32
hypothetical protein FLJ20307	NM_017757	1.32
golgi associated, gamma adaptin ear containing, ARF binding protein 3	AF219138	1.32
SRY (sex determining region Y)-box 20	NM_006942	1.32
Fanconi anemia, complementation group G	AC004472	1.32
Microfibril-associated glycoprotein-2	U37283	1.32
TERA protein	NM_021238	1.32
hypothetical protein FLJ12270	NM_030581	1.32
gb:NM_004250.1 /DEF=Homo sapiens RAB9, member RAS oncogene family, pseudogene 1 (RAB9P1), mRNA. /FEA=mRNA /GEN=RAB9P1 /PROD=RAB9, member RAS oncogene family, pse	NM_004250	1.32
Consensus includes gb:AL049301.1 /DEF=Homo sapiens mRNA; cDNA DKFZp564P073 (from clone DKFZp564P073). /FEA=mRNA /DB_XREF=gi:4500067 /UG=Hs.278577 Homo sapiens mI	AL049301	1.32
CGI-79 protein	AF078865	1.32
caspase 8, apoptosis-related cysteine protease	NM_001228	1.32
stress 70 protein chaperone, microsome-associated, 60kD	AJ718418	1.32
smoothened homolog (Drosophila)	NM_005631	1.32
solute carrier family 22 (organic anion transporter), member 7	AF210455	1.32
tumor necrosis factor (ligand) superfamily, member 18	NM_005092	1.32
complement component 1, q subcomponent binding protein	BE964082	1.32
procollagen-lysine, 2-oxoglutarate 5-dioxygenase (lysine hydroxylase, Ehlers-Danlos syndrome type VI)	NM_000302	1.32
N-acetylglucosamine-1-phosphodiester alpha-N-acetylglucosaminidase	NM_016256	1.32
zinc finger protein 295	AP001745	1.32
sialyltransferase 1 (beta-galactoside alpha-2,6-sialyltransferase)	X17247	1.32
CGI-142	AB029156	1.32
hypothetical protein FLJ10846	NM_018241	1.32
mannosidase, alpha, class 2C, member 1	NM_006715	1.32
zinc finger protein 14 (KOX 6)	NM_021030	1.32
alkaline phosphatase, placental (Regan isozyme)	M13077	1.32
oculocutaneous albinism II (pink-eye dilution (murine) homolog)	NM_000275	1.32
cadherin, EGF LAG seven-pass G-type receptor 1 (flamingo homolog, Drosophila)	AW241864	1.32
zinc finger protein 202	N46430	1.32
Consensus includes gb:BF344237 /FEA=EST /DB_XREF=gi:11291450 /DB_XREF=est:602017491F1 /CLONE=IMAGE:4152983 /UG=Hs.124776 Homo sapiens mRNA; cDNA DKFZp564N111	BF344237	1.32
hypothetical protein FLJ11088	NM_018318	1.32
MAD, mothers against decapentaplegic homolog 6 (Drosophila)	NM_005585	1.32
Consensus includes gb:AF184174 /DEF=Homo sapiens somatostatin receptor 2 (SSTR2) gene, complete cds, alternatively spliced /FEA=CDS_2 /DB_XREF=gi:7229402 /UG=Hs.184841 son	AF184174	1.32
RAB26, member RAS oncogene family	NM_014353	1.32
gap junction protein, alpha 4, 37kD (connexin 37)	NM_002060	1.32
gb:NM_014823.1 /DEF=Homo sapiens KIAA0344 gene product (KIAA0344), mRNA. /FEA=mRNA /GEN=KIAA0344 /PROD=KIAA0344 gene product /DB_XREF=gi:7662065 /UG=Hs.321444 I	NM_014823	1.32
UL16 binding protein 2	NM_025217	1.32
hypothetical protein FLJ11274	NM_018375	1.32
tRNA isopentenylpyrophosphate transferase	AF116679	1.32

KIAA0671 gene product	NM_014011	1.32
desmoplakin (DPI, DPII)	NM_004415	1.32
growth hormone releasing hormone	NM_021081	1.32
Consensus includes gb:AK023774.1 /DEF=Homo sapiens cDNA FLJ13712 fis, clone PLACE2000394. /FEA=mRNA /DB_XREF=gi:10435808 /UG=Hs.273099 Homo sapiens cDNA FLJ13712 fi	AK023774	1.32
adenylate cyclase 9	NM_001116	1.32
protein kinase, lysine deficient 1	AI768512	1.32
Consensus includes gb:AL049593 /DEF=Human DNA sequence from clone RP4-654A7 on chromosome 20. Contains the gene for phospholipase C-beta-1 (KIAA0581, 1-phosphatidylinositol-	AL049593	1.32
adenosine deaminase, tRNA-specific 1	NM_012091	1.32
kinesin family member 3C	NM_002254	1.32
deoxyribonuclease I-like 1	NM_006730	1.32
potassium channel, subfamily K, member 2 (TREK-1)	AF004711	1.32
ALEX1 protein	NM_016608	1.32
alpha-2-macroglobulin	NM_000014	1.32
cholecystokinin	NM_000729	1.32
amyloid beta (A4) precursor protein (protease nexin-II, Alzheimer disease)	BC004369	1.32
adenylate kinase 2	AW277253	1.32
homologous to yeast nitrogen permease (candidate tumor suppressor)	AC002481	1.32
MOST2 protein	NM_020250	1.32
G protein-coupled receptor, family C, group 5, member B	AF202640	1.32
melanoma antigen, family B, 4	NM_002367	1.32
hypothetical protein FLJ10637	AF274950	1.32
hypothetical protein FLJ10376	NM_018076	1.32
lipocalin 2 (oncogene 24p3)	NM_005564	1.32
luteinizing hormone/choriogonadotropin receptor	NM_000233	1.32
pleckstrin homology-like domain, family A, member 1	NM_007350	1.32
DKFZP547G910 protein	AL049925	1.32
p53-induced protein PIGPC1	NM_022121	1.32
dipeptidylpeptidase VI	NM_001936	1.32
glioma-amplified sequence-41	NM_006530	1.32
RE1-silencing transcription factor	AI978576	1.32
EHM2 gene	NM_019114	1.32
small inducible cytokine subfamily D (Cys-X3-Cys), member 1 (fractalkine, neurotactin)	U84487	1.32
KIAA0130 gene product	AK022508	1.32
1-acylglycerol-3-phosphate O-acyltransferase 1 (lysophosphatidic acid acyltransferase, alpha)	AF007145	1.32
karyopherin alpha 6 (importin alpha 7)	BG403834	1.32
Consensus includes gb:AK021614.1 /DEF=Homo sapiens cDNA FLJ11552 fis, clone HEMBA1003021. /FEA=mRNA /DB_XREF=gi:10432830 /UG=Hs.306606 Homo sapiens cDNA FLJ11552 f	AK021614	1.32
alkaline phosphatase, placental-like 2	J04948	1.32
Consensus includes gb:AI249275 /DEF=Homo sapiens partial MTHFR gene for methylenetetrahydrofolate reductase /FEA=CDS /DB_XREF=gi:7271187 /UG=Hs.214142 5,10-methylenetetra	AI249275	1.32
Charot-Leyden crystal protein	NM_001828	1.32
BarH-like homeobox 2	AF031924	1.32
KIAA0665 gene product	AW243089	1.32
syndecan 1	Z48199	1.32
microtubule-associated protein 2	U89330	1.32
Consensus includes gb:AL049259.1 /DEF=Homo sapiens mRNA; cDNA DKFZp564E193 (from clone DKFZp564E193). /FEA=mRNA /DB_XREF=gi:4500005 /UG=Hs.11056 Homo sapiens mR	AL049259	1.32
CDC14 cell division cycle 14 homolog A (S. cerevisiae)	AF064103	1.32
Wilms' tumour 1-associating protein	AL583911	1.32
WD repeat domain 7	AB011113	1.32
DKFZP434N014 protein	BE045998	1.32
ankyrin 1, erythrocytic	NM_020480	1.32
C3HC4-like zinc finger protein	NM_016422	1.32
protein phosphatase 1A (formerly 2C), magnesium-dependent, alpha isoform	AF070670	1.32
mesoderm homolog 1 (Drosophila)	AC003982	1.32
Consensus includes gb:AL109681.1 /DEF=Homo sapiens mRNA full length insert cDNA clone EUROIMAGE 112333. /FEA=mRNA /DB_XREF=gi:5689785 /UG=Hs.226017 Homo sapiens mRN	AL109681	1.32
ATP synthase, H+ transporting, mitochondrial F0 complex, subunit g	AA917672	1.32
phosphoribosyl pyrophosphate amidotransferase	AI457120	1.32
hemochromatosis	BG402460	1.32
Consensus includes gb:AI921841 /FEA=EST /DB_XREF=gi:5657805 /DB_XREF=est:wp07a09.x1 /CLONE=IMAGE:2464120 /UG=Hs.212670 ESTs	AI921841	1.32
Consensus includes gb:AU143984 /FEA=EST /DB_XREF=gi:11005505 /DB_XREF=est:AU143984 /CLONE=HEMBA1000568 /UG=Hs.301645 Homo sapiens cDNA FLJ10021 fis, clone HEMBA	AU143984	1.32
Kruppel-like factor 8	NM_007250	1.32
Consensus includes gb:AK024177.1 /DEF=Homo sapiens cDNA FLJ14115 fis, clone MAMMA1001760. /FEA=mRNA /DB_XREF=gi:10436493 /UG=Hs.266263 Homo sapiens cDNA FLJ14115 f	AK024177	1.32
v-jun sarcoma virus 17 oncogene homolog (avian)	BC002646	1.32
RAN binding protein 3	BC004349	1.32
visinin-like 1	AF039555	1.32

opioid growth factor receptor	AF172450	1.32
stathmin-like 2	NM_007029	1.32
hypothetical protein MGC2592	AL121891	1.32
ficolin (collagen/fibrinogen domain-containing) 1	NM_002003	1.32
nuclear receptor subfamily 2, group F, member 2	AV703465	1.32
phospholipase A2, group IVA (cytosolic, calcium-dependent)	M68874	1.32
solute carrier family 19 (folate transporter), member 1	BC003068	1.32
Consensus includes gb:AV733950 /FEA=EST /DB_XREF=gi:10851495 /DB_XREF=est:AV733950 /CLONE=cdAAD612 /UG=Hs.326035 early growth response 1 /FL=gb:M62829.1 gb:NM_0(	AV733950	1.32
aldehyde dehydrogenase 1 family, member A3	NM_000693	1.32
prefoldin 4	AI718295	1.32
CGG triplet repeat binding protein 1	AI694317	1.32
zinc finger protein 135 (clone pHZ-17)	NM_003436	1.32
hypothetical protein	AC002550	1.32
glycine dehydrogenase (decarboxylating; glycine decarboxylase, glycine cleavage system protein P)	NM_000170	1.32
lamin A/C	AK026584	1.32
microphthalmia-associated transcription factor	NM_000248	1.32
hypothetical protein FLJ11110	NM_018326	1.32
sialophorin (gpL115, leukosialin, CD43)	X52075	1.32
hypothetical protein FLJ20014	NM_017622	1.32
anti-Mullerian hormone receptor, type II	NM_020547	1.32
neurexophilin 4	AI933199	1.32
tumor necrosis factor receptor superfamily, member 6	Z70519	1.32
regulatory factor X, 3 (influences HLA class II expression)	NM_002919	1.32
leukocyte membrane antigen	AF161346	1.32
calreticulin	AD000092	1.32
Consensus includes gb:AL137428.1 /DEF=Homo sapiens mRNA; cDNA DKFZp761N1323 (from clone DKFZp761N1323). /FEA=mRNA /DB_XREF=gi:6807985 /UG=Hs.306459 Homo sapiens	AL137428	1.32
KIAA1045 protein	AB028968	1.32
Consensus includes gb:AF339785.1 /DEF=Homo sapiens clone IMAGE:1963178, mRNA sequence. /FEA=mRNA /DB_XREF=gi:13507323 /UG=Hs.270256 Homo sapiens clone IMAGE:19631;	AF339785	1.32
protein-O-mannosyltransferase 1	NM_007171	1.32
Consensus includes gb:BE045982 /FEA=EST /DB_XREF=gi:8363035 /DB_XREF=est:hd91b04.x4 /CLONE=IMAGE:2916847 /UG=Hs.99093 Homo sapiens chromosome 19, cosmid R28379	BE045982	1.32
macrophage scavenger receptor 1	NM_002445	1.32
chromodomain helicase DNA binding protein 3	BE379542	1.32
cyfticin, basic protein of sperm head cytoskeleton 1	Z22780	1.32
hypothetical protein MGC2470	NM_024331	1.32
KIAA0403 protein	AW166711	1.32
gb:AF116659.1 /DEF=Homo sapiens PRO1412 mRNA, complete cds. /FEA=mRNA /PROD=PRO1412 /DB_XREF=gi:7959816 /UG=Hs.321151 Homo sapiens PRO1412 mRNA, complete cds /	AF116659	1.32
Consensus includes gb:AK026679.1 /DEF=Homo sapiens cDNA: FLJ23026 fis, clone LNG01738. /FEA=mRNA /DB_XREF=gi:10439585 /UG=Hs.306863 Homo sapiens cDNA: FLJ23026 fis, cl	AK026679	1.32
nucleosome assembly protein 1-like 2	NM_021963	1.32
Consensus includes gb:AK000185.1 /DEF=Homo sapiens cDNA FLJ20178 fis, clone COL09990. /FEA=mRNA /DB_XREF=gi:7020104 /UG=Hs.306389 Homo sapiens cDNA FLJ20178 fis, clon	AK000185	1.32
diaphanous homolog 2 (Drosophila)	NM_006729	1.32
retinal short-chain dehydrogenase/reductase retSDR3	NM_016246	1.32
killer cell immunoglobulin-like receptor, two domains, short cytoplasmic tail, 2	L76668	1.32
myosin regulatory light chain	U26162	1.32
potassium voltage-gated channel, subfamily H (eag-related), member 4	NM_012285	1.32
hypothetical protein FLJ21277	NM_025033	1.32
calcium channel, voltage-dependent, alpha 1I subunit	AB032946	1.32
gb:BC006408.1 /DEF=Homo sapiens, Similar to zinc finger protein 254, clone MGC:10544, mRNA, complete cds. /FEA=mRNA /PROD=Similar to zinc finger protein 254 /DB_XREF=gi:13623	BC006408	1.32
phosphatidylinositol glycan, class A (paroxysmal nocturnal hemoglobinuria)	NM_002641	1.32
Consensus includes gb:AW975050 /FEA=EST /DB_XREF=gi:8166254 /DB_XREF=est:EST387155 /UG=Hs.293892 ESTs, Weakly similar to ALU4_HUMAN ALU SUBFAMILY SB2 SEQUENCE CON	AW975050	1.32
Consensus includes gb:AL137457.1 /DEF=Homo sapiens mRNA; cDNA DKFZp566G231 (from clone DKFZp566G231). /FEA=mRNA /DB_XREF=gi:6808035 /UG=Hs.272324 Homo sapiens mI	AL137457	1.32
cytokine receptor-like factor 1	NM_004750	1.32
uroplakin 1B	AB002155	1.32
N-acetylgalactosaminidase, alpha-	NM_000262	1.32
putative integral membrane transporter	T15777	1.32
KIAA0453 protein	AA524553	1.32
N-deacetylase/N-sulfotransferase 4	NM_022569	1.32
Consensus includes gb:AI744673 /FEA=EST /DB_XREF=gi:5112883 /DB_XREF=est:wg02f04.x1 /CLONE=IMAGE:2363935 /UG=Hs.186970 ESTs	AI744673	1.32
neurotrimin	AW117456	1.32
interleukin 18 binding protein	NM_005699	1.32
olfactory receptor, family 2, subfamily F, member 1	NM_012369	1.32
hypothetical protein FLJ13955	NM_024759	1.32
frizzled-related protein	NM_001463	1.32
Consensus includes gb:AU145890 /FEA=EST /DB_XREF=gi:11007411 /DB_XREF=est:AU145890 /CLONE=HEMBA1006158 /UG=Hs.284186 forkhead box C1 /FL=gb:NM_001453.1	AU145890	1.32

hypothetical protein MGC16714	BE792298	1.32
UDP-glucose ceramide glucosyltransferase-like 1	NM_020120	1.32
X-ray repair complementing defective repair in Chinese hamster cells 4	BC005259	1.32
UDP-N-acetyl-alpha-D-galactosamine:(N-acetylneuraminyl)-galactosylglucosylceramide N-acetylgalactosaminyltransferase (GalNAc-T)	NM_001478	1.32
catenin (cadherin-associated protein), alpha 1 (102kD)	D14705	1.32
likely homolog of mouse heparin/heparin sulfate-glucuronic acid C5-epimerase	W87398	1.32
LIM homeobox protein 5	NM_022363	1.32
triple functional domain (PTPRF interacting)	AI797657	1.32
amine oxidase, copper containing 2 (retina-specific)	NM_009590	1.32
supervillin	AF051851	1.31
pre-B-cell leukemia transcription factor 2	NM_002586	1.31
Cluster Incl. AI743990:wc32d07.x1 Homo sapiens cDNA, 3' end /clone=IMAGE-2316877 /clone_end=3' /gb=AI743990 /gi=5112278 /ug=Hs.24859 /len=583	AI743990	1.31
pleckstrin homology-like domain, family A, member 1	NM_007350	1.31
Toll-like receptor 8	NM_016610	1.31
Consensus includes gb:U79249.1 /DEF=Human clone 23839 mRNA sequence. /FEA=mRNA /DB_XREF=gi:1710196 /UG=Hs.78362 Human clone 23839 mRNA sequence	U79249	1.31
parathyroid hormone	NM_000315	1.31
Consensus includes gb:AW971205 /FEA=EST /DB_XREF=gi:8161050 /DB_XREF=est:EST383293 /UG=Hs.114280 ESTs	AW971205	1.31
Consensus includes gb:AF070591.1 /DEF=Homo sapiens clone 24649 mRNA sequence. /FEA=mRNA /DB_XREF=gi:3387966 /UG=Hs.322473 Homo sapiens clone 24649 mRNA sequence	AF070591	1.31
absent in melanoma 2	NM_004833	1.31
fatty-acid-Coenzyme A ligase, long-chain 6	AV727634	1.31
phosphatidylinositol 4-kinase type II	BC003167	1.31
immunoglobulin superfamily, member 3	AB007935	1.31
interferon induced transmembrane protein 2 (1-8D)	NM_006435	1.31
aconitase 2, mitochondrial	AL021877	1.31
seizure related 6 homolog (mouse)-like	AL050253	1.31
hypothetical protein FLJ23191	NM_024574	1.31
gap junction protein, beta 3, 31kD (connexin 31)	BF060667	1.31
neurofilament, heavy polypeptide (200kD)	X15306	1.31
basic transcription factor 3	R83000	1.31
RAN binding protein 3	NM_007321	1.31
fusion, derived from t(12;16) malignant liposarcoma	AW514140	1.31
hypothetical protein FLJ23058	NM_024696	1.31
defensin, alpha 1, myeloid-related sequence	NM_004084	1.31
H1 histone family, member 3	AL031777	1.31
protein tyrosine phosphatase, receptor type, K	NM_002844	1.31
transmembrane protease, serine 2	NM_005656	1.31
glucose-6-phosphatase, catalytic (glycogen storage disease type I, von Gierke disease)	NM_000151	1.31
caspase 6, apoptosis-related cysteine protease	U20537	1.31
pyruvate carboxylase	NM_022172	1.31
SEC24 related gene family, member D (S. cerevisiae)	NM_014822	1.31
cytosolic ovarian carcinoma antigen 1	AF207881	1.31
DKFZP547G0910 protein	AL049925	1.31
RNA-binding protein gene with multiple splicing	D84109	1.31
protocadherin 11	AF332218	1.31
TATA box binding protein (TBP)-associated factor, RNA polymerase I, B, 63kD	AA004579	1.31
TNFRSF1A-associated via death domain	NM_003789	1.31
Consensus includes gb:AK023187.1 /DEF=Homo sapiens cDNA FLJ13125 fis, clone NT2RP3002877. /FEA=mRNA /DB_XREF=gi:10435001 /UG=Hs.287551 Homo sapiens cDNA FLJ13125 fi	AK023187	1.31
programmed cell death 4 (neoplastic transformation inhibitor)	NM_014456	1.31
T-box 1	AF012130	1.31
Consensus includes gb:AL049301.1 /DEF=Homo sapiens mRNA; cDNA DKFZp564P073 (from clone DKFZp564P073). /FEA=mRNA /DB_XREF=gi:4500067 /UG=Hs.278577 Homo sapiens mI	AL049301	1.31
one cut domain, family member 2	NM_004852	1.31
GRO1 oncogene (melanoma growth stimulating activity, alpha)	NM_001511	1.31
DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 28	BG477502	1.31
peanut-like 1 (Drosophila)	AA702163	1.31
calmodulin-like 3	M36707	1.31
gene for serine/threonine protein kinase	NM_018401	1.31
neurofilament, light polypeptide (68kD)	AL537457	1.31
cyclin-dependent kinase inhibitor 1C (p57, Kip2)	R78668	1.31
Consensus includes gb:AC003989 /DEF=Human BAC clone CTB-7J15 from 7q31 /FEA=CDS /DB_XREF=gi:2772538 /UG=Hs.248069 Human BAC clone CTB-7J15 from 7q31	AC003989	1.31
KIAA1100 protein	NM_014901	1.31
histone deacetylase 6	AK024083	1.31
discs, large (Drosophila) homolog-associated protein 2	AF009204	1.31
delta-tubulin	NM_016261	1.31

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aryl hydrocarbon receptor nuclear translocator-like	AB000812	1.31
matrix metalloproteinase 9 (gelatinase B, 92kD gelatinase, 92kD type IV collagenase)	AL162458	1.31
mucin 3A, intestinal	AF113616	1.31
dual specificity phosphatase 10	N36770	1.31
SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily c, member 2	NM_003075	1.31
Consensus includes gb:AL137547.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434B1120 (from clone DKFZp434B1120). /FEA=mRNA /DB_XREF=gi:6808230 /UG=Hs.259619 Homo sapiens	AL137547	1.31
collagen, type IV, alpha 1	NM_001845	1.31
pleiomorphic adenoma gene-like 1	NM_002656	1.31
G protein-coupled receptor 42	NM_005305	1.31
zinc finger protein 306	BC006118	1.31
gb:AF130057.1 /DEF=Homo sapiens clone FLB5539 PRO1454 mRNA, complete cds. /FEA=mRNA /PROD=PRO1454 /DB_XREF=gi:11493420 /UG=Hs.323288 Homo sapiens clone FLB5539	AF130057	1.31
sulfotransferase family, cytosolic, 1C, member 1	AF186255	1.31
RAB31, member RAS oncogene family	NM_006868	1.31
decay accelerating factor for complement (CD55, Cromer blood group system)	NM_000574	1.31
phosphoserine phosphatase	NM_004577	1.31
ceroid-lipofuscinosis, neuronal 3, juvenile (Batten, Spielmeyer-Vogt disease)	AC002544	1.31
deleted in azoospermia	U21663	1.31
Consensus includes gb:AI590053 /FEA=EST /DB_XREF=gi:4599101 /DB_XREF=est:tr75d05.x1 /CLONE=IMAGE:2224137 /UG=Hs.235160 Homo sapiens mRNA; cDNA DKFZp434G0572 (fro	AI590053	1.31
Consensus includes gb:AL050032.1 /DEF=Homo sapiens mRNA; cDNA DKFZp566F1224 (from clone DKFZp566F1224). /FEA=mRNA /DB_XREF=gi:4884272 /UG=Hs.306307 Homo sapiens	AL050032	1.31
rab3 interacting protein	AF263310	1.31
prostate epithelium-specific Ets transcription factor	AI745526	1.31
ataxia telangiectasia mutated (includes complementation groups A, C and D)	U26455	1.31
hypothetical protein FLJ22170	NM_025099	1.31
ret proto-oncogene (multiple endocrine neoplasia and medullary thyroid carcinoma 1, Hirschsprung disease)	X15786	1.31
hypothetical protein MGC5590	NM_024058	1.31
ectonucleotide pyrophosphatase/phosphodiesterase 4 (putative function)	NM_014936	1.31
peroxisome proliferative activated receptor, alpha	NM_005036	1.31
pellino homolog 2 (Drosophila)	NM_021255	1.31
mitogen-activated protein kinase 13	BC000433	1.31
MADS box transcription enhancer factor 2, polypeptide C (myocyte enhancer factor 2C)	NM_002397	1.31
fibrillin 1 (Marfan syndrome)	NM_000138	1.31
calponin like transmembrane domain protein	NM_024734	1.31
hypothetical protein FLJ13150	NM_024813	1.31
protocadherin alpha 12	AF152308	1.31
solute carrier family 13 (sodium-dependent dicarboxylate transporter), member 2	NM_003984	1.31
MDS024 protein	NM_021820	1.31
heart and neural crest derivatives expressed 1	NM_004821	1.31
ankyrin repeat domain 6	NM_014942	1.31
MHC class I region ORF	NM_006674	1.31
disrupted in schizophrenia 1	NM_018662	1.31
ATPase, Class II, type 9B	R50094	1.31
hypothetical protein FLJ13687	NM_024935	1.31
orphan seven-transmembrane receptor, chemokine related	NM_016557	1.31
Consensus includes gb:AL133571.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434F1135 (from clone DKFZp434F1135). /FEA=mRNA /DB_XREF=gi:6599149 /UG=Hs.116128 Homo sapiens	AL133571	1.31
isoprenylcysteine carboxyl methyltransferase	AF064084	1.31
serine protease inhibitor-like, with Kunitz and WAP domains 1 (eppin)	AF286368	1.31
Consensus includes gb:AL139112 /DEF=Human DNA sequence from clone GS1-103B18 on chromosome Xq27.1-27.3 Contains ESTs, STSs and GSSs. Contains a novel cytochrome c-1 (CYC1)	AL139112	1.31
TATA box binding protein (TBP)-associated factor, RNA polymerase I, C, 110kD	NM_005679	1.31
estrogen receptor 1	NM_000125	1.31
killer cell immunoglobulin-like receptor, three domains, long cytoplasmic tail, 1	AF262973	1.31
talin 2	AV728958	1.31
osteomodulin	AI765819	1.31
hypothetical protein FLJ10811	NM_018228	1.31
Consensus includes gb:AL049710 /DEF=Human DNA sequence from clone 633H17 on chromosome 1p31.2-32.2. Contains a pseudogene similar to part of MTCO1 (Cytochrome C oxidase 1).	AL049710	1.31
colony stimulating factor 1 (macrophage)	U22386	1.31
hypothetical protein FLJ21125	NM_024627	1.31
Consensus includes gb:AI188346 /FEA=EST /DB_XREF=gi:3739555 /DB_XREF=est:qd08a10.x1 /CLONE=IMAGE:1723098 /UG=Hs.301776 ESTs, Weakly similar to ALU1_HUMAN ALU SUBFA	AI188346	1.31
solute carrier family 17 (sodium-dependent inorganic phosphate cotransporter), member 6	NM_020346	1.31
chitobiase, di-N-acetyl-	AW304174	1.31
solute carrier family 11 (proton-coupled divalent metal ion transporters), member 2	AF064484	1.31
adrenergic, alpha-2A-, receptor	AF284095	1.31
HYA22 protein	AA129726	1.31
66C protein	NM_025261	1.31

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myeloid leukemia factor 1	NM_022443	1.31
phospholipase D1, phosphatidylcholine-specific	NM_002662	1.31
U2 small nuclear ribonucleoprotein auxiliary factor, small subunit 2	AW089584	1.31
dentatorubral-pallidoluysian atrophy (atrophin-1)	U47924	1.31
B9 protein	BC002944	1.31
amiloride-sensitive cation channel 2, neuronal	NM_020039	1.31
holocytochrome c synthase (cytochrome c heme-lyase)	AI801013	1.31
crystallin, gamma C	NM_020989	1.31
DKFZP547E2110 protein	BF062335	1.31
zinc finger protein 26 (KOX 20)	NM_019591	1.31
protein expressed in thyroid	NM_014297	1.31
Consensus includes gb:AI378044 /FEA=EST /DB_XREF=gi:4187897 /DB_XREF=est:te67h09.x1 /CLONE=IMAGE:2091809 /UG=Hs.23703 ESTs	AI378044	1.31
hook1 protein	NM_015888	1.31
Consensus includes gb:AU144792 /FEA=EST /DB_XREF=gi:11006313 /DB_XREF=est:AU144792 /CLONE=HEMBA1002973 /UG=Hs.319540 Homo sapiens cDNA FLJ10127 fis, clone HEMBA	AU144792	1.31
HSKM-B protein	AI809870	1.31
ELK4, ETS-domain protein (SRF accessory protein 1)	NM_021795	1.31
Ral guanine nucleotide exchange factor RalGPS1A	NM_014636	1.31
Rho guanine exchange factor (GEF) 16	NM_014448	1.31
myeloid/lymphoid or mixed-lineage leukemia (trithorax homolog, Drosophila)	AI701430	1.31
solute carrier family 7 (cationic amino acid transporter, y+ system), member 1	NM_003045	1.31
hypothetical protein FLJ21347	NM_022827	1.31
secreted protein of unknown function	AI984136	1.31
interferon, alpha 8	NM_002170	1.31
T cell receptor gamma constant 2	AL580863	1.31
zinc finger protein, subfamily 1A, 2 (Helios)	NM_016260	1.31
proline arginine-rich end leucine-rich repeat protein	NM_002725	1.31
phosphate cytidylyltransferase 1, choline, alpha isoform	AI638771	1.31
mesoshl homolog 1 (Drosophila)	NM_002442	1.31
solute carrier family 37 (glycerol-3-phosphate transporter), member 1	NM_018964	1.31
SRY (sex determining region Y)-box 10	AI367319	1.31
hypothetical protein FLJ20516	NM_017858	1.31
HIV-1 Rev binding protein	NM_004504	1.31
cholecystokinin B receptor	BC000740	1.31
Consensus includes gb:BE045142 /FEA=EST /DB_XREF=gi:8362195 /DB_XREF=est:hn26h02.x1 /CLONE=IMAGE:3023283 /UG=Hs.274907 ESTs	BE045142	1.31
Consensus includes gb:BG434174 /FEA=EST /DB_XREF=gi:13340680 /DB_XREF=est:602506106F1 /CLONE=IMAGE:4603513 /UG=Hs.23437 Homo sapiens cDNA FLJ13555 fis, clone PLAC	BG434174	1.31
KIAA1031 protein	NM_025013	1.31
KIAA0962 protein	AB023179	1.31
hypothetical protein FLJ20920	NM_025149	1.31
retinoblastoma binding protein 4	AL390237	1.31
hypothetical protein FLJ23259	NM_024727	1.31
melanoma antigen, family A, 8	BC002455	1.31
KIAA0535 gene product	NM_014682	1.31
protein-kinase, interferon-inducible double stranded RNA dependent inhibitor, repressor of (P58 repressor)	AF081567	1.31
lipase, hormone-sensitive	NM_005357	1.31
hypothetical protein FLJ10895	NM_019084	1.31
Consensus includes gb:AK024527.1 /DEF=Homo sapiens cDNA: FLJ20874 fis, clone ADKA02818. /FEA=mRNA /DB_XREF=gi:10436829 /UG=Hs.306684 Homo sapiens cDNA: FLJ20874 fis,	AK024527	1.31
KIAA0557 protein	BF508774	1.31
collagen, type IX, alpha 1	NM_001851	1.31
paraoxonase 1	U53784	1.31
olfactory receptor, family 2, subfamily H, member 1	AI302604	1.31
Consensus includes gb:AK025580.1 /DEF=Homo sapiens cDNA: FLJ21927 fis, clone HEP04178, highly similar to HSU90909 Human clone 23722 mRNA sequence. /FEA=mRNA /DB_XREF=g	AK025580	1.31
KIAA0848 protein	NM_014926	1.31
hypothetical protein FLJ13892	NM_025017	1.31
thyroid stimulating hormone receptor	BE740743	1.31
Mad4 homolog	BC002713	1.31
protocadherin gamma subfamily A, 3	AF152510	1.31
mitogen-activated protein kinase kinase kinase 12	NM_006301	1.31
triosephosphate isomerase 1	M10036	1.31
glycine receptor, beta	NM_000824	1.31
trinucleotide repeat containing 11 (THR-associated protein, 230 kDa subunit)	AI217362	1.31
Consensus includes gb:AL049690 /DEF=Human DNA sequence from clone 1060K6 on chromosome 20p12.1-13 Contains a pseudogene similar to 40S ribosomal protein S11, ESTs, STSs and	AL049690	1.31
Retina-derived POU-domain factor-1	NM_007252	1.31
Consensus includes gb:X61072.1 /DEF=Human mRNA for T cell receptor, clone IGRA17. /FEA=mRNA /PROD=T cell receptor /DB_XREF=gi:33511 /UG=Hs.272500 Human mRNA for T cell re	X61072	1.31

sodium channel, voltage-gated, type II, beta polypeptide	AF107028	1.31
Friedreich ataxia region gene X123	AA524029	1.31
KIAA0336 gene product	NM_014635	1.31
hypothetical protein FLJ21522	NM_024526	1.31
CD84 antigen (leukocyte antigen)	U96627	1.31
sialyltransferase 4C (beta-galactosidase alpha-2,3-sialytransferase)	NM_006278	1.31
zinc finger protein 41	AI927984	1.31
nuclear receptor co-repressor 2	S83390	1.31
matrix metalloproteinase 26	NM_021801	1.31
phosphodiesterase 9A	NM_002606	1.31
carcinoembryonic antigen-related cell adhesion molecule 1 (biliary glycoprotein)	M69176	1.31
hypothetical protein FLJ20489	H14241	1.30
MAX protein	NM_002382	1.30
WW domain-containing protein 1	AU155187	1.30
ATPase, H+/K+ exchanging, alpha polypeptide	NM_000704	1.30
regulator of G-protein signalling 4	AL514445	1.30
gb:BC000453.1 /DEF=Homo sapiens, Similar to pericentriolar material 1, clone MGC:8458, mRNA, complete cds. /FEA=mRNA /PROD=Similar to pericentriolar material 1 /DB_XREF=gi:1265	BC000453	1.30
KIAA1032 protein	BE783668	1.30
sperm associated antigen 11	NM_016512	1.30
retinitis pigmentosa GTPase regulator interacting protein 1	NM_020366	1.30
olfactory receptor, family 1, subfamily D, member 2	U53583	1.30
protocadherin 16	NM_024542	1.30
CD86 antigen (CD28 antigen ligand 2, B7-2 antigen)	L25259	1.30
keratin 13	NM_002274	1.30
RAD21 homolog (S. pombe)	NM_006265	1.30
T cell receptor beta locus	AF009787	1.30
histidine-rich glycoprotein	NM_000412	1.30
ocular albinism 1 (Nettleship-Falls)	NM_000273	1.30
mitogen-activated protein kinase-activated protein kinase 5	NM_003668	1.30
Consensus includes gb:AL080082.1 /DEF=Homo sapiens mRNA; cDNA DKFZp56461162 (from clone DKFZp56461162). /FEA=mRNA /DB_XREF=gi:5262495 /UG=Hs.139006 Homo sapiens	AL080082	1.30
splicing factor, arginine/serine-rich 2, interacting protein	AW084759	1.30
KIAA0924 protein	NM_014897	1.30
glutamate receptor, metabotropic 4	U92457	1.30
cyclin-dependent kinase inhibitor 2D (p19, inhibits CDK4)	U20498	1.30
solute carrier family 7 (cationic amino acid transporter, y+ system), member 4	NM_004173	1.30
eukaryotic translation initiation factor 4 gamma, 3	NM_003760	1.30
olfactory receptor, family 7, subfamily E, member 24 pseudogene	BF062364	1.30
histone deacetylase 6	BC005872	1.30
KIAA1043 protein	AB028966	1.30
trefoil factor 2 (spasmolytic protein 1)	NM_005423	1.30
iduronate 2-sulfatase (Hunter syndrome)	AI819115	1.30
breast carcinoma amplified sequence 1	NM_003657	1.30
transglutaminase 4 (prostate)	NM_003241	1.30
Consensus includes gb:AK022094.1 /DEF=Homo sapiens cDNA FLJ12032 fis, clone HEMBB1001880. /FEA=mRNA /DB_XREF=gi:10433415 /UG=Hs.151194 Homo sapiens cDNA FLJ12032 f	AK022094	1.30
KIAA0339 gene product	N30342	1.30
asialoglycoprotein receptor 2	NM_001181	1.30
tocopherol (alpha) transfer protein (ataxia (Friedreich-like) with vitamin E deficiency)	U21938	1.30
Consensus includes gb:AK001155.1 /DEF=Homo sapiens cDNA FLJ10293 fis, clone NT2RM1000280, highly similar to VACUOLAR ATP SYNTHASE SUBUNIT D (EC 3.6.1.34). /FEA=mRNA /DB_	AK001155	1.30
glutathione S-transferase M2 (muscle)	NM_000848	1.30
hypothetical protein FLJ13544	NM_025008	1.30
hypothetical protein FLJ10210	NM_018027	1.30
endosulfine alpha	AF067170	1.30
activator of CREM in testis	NM_020482	1.30
neuropeptide FF 1; RFamide-related peptide receptor	NM_022146	1.30
Consensus includes gb:AK026484.1 /DEF=Homo sapiens cDNA: FLJ22831 fis, clone KAIA4161. /FEA=mRNA /DB_XREF=gi:10439356 /UG=Hs.321666 Homo sapiens cDNA: FLJ22831 fis, cl	AK026484	1.30
hypothetical protein	AI740515	1.30
collagen, type XXI, alpha 1	NM_030820	1.30
protocadherin 1 (cadherin-like 1)	NM_002587	1.30
protein phosphatase 4 regulatory subunit 2	NM_019853	1.30
talin 2	AB002318	1.30
hypothetical protein FLJ20527	NM_017863	1.30
biotinidase	AI767414	1.30
glycogenin 2	U94363	1.30

neuropilin 1	AF145712	1.30
chromosome 1 open reading frame 15	AF288395	1.30
MAS1 oncogene	NM_002377	1.30
hypothetical protein FLJ23468	NM_024629	1.30
Langerhans cell specific c-type lectin	NM_015717	1.30
thrombospondin 1	BE999967	1.30
cellular repressor of E1A-stimulated genes	NM_003851	1.30
centaurin, alpha 1	NM_006869	1.30
alpha-2-HS-glycoprotein	NM_001622	1.30
small inducible cytokine subfamily A (Cys-Cys), member 17	NM_002987	1.30
neurotensin	NM_006183	1.30
properdin P factor, complement	NM_002621	1.30
tripartite motif-containing 15	U34249	1.30
solute carrier family 12 (potassium/chloride transporters), member 4	AF047338	1.30
carboxyl ester lipase (bile salt-stimulated lipase)	NM_001807	1.30
Consensus includes gb:AI890903 /FEA=EST /DB_XREF=gi:5596067 /DB_XREF=est:wm91i10.x1 /CLONE=IMAGE:2443339 /UG=Hs.119683 ESTs	AI890903	1.30
chondroitin sulfate proteoglycan 5 (neuroglycan C)	NM_006574	1.30
hypothetical protein FLJ23138	AI743090	1.30
ubiquitin-conjugating enzyme E2I (UBC9 homolog, yeast)	AA910614	1.30
interferon, alpha 14	NM_002172	1.30
hypothetical protein SB8167	NM_020393	1.30
E74-like factor 5 (ets domain transcription factor)	AF115403	1.30
matrix metalloproteinase 24 (membrane-inserted)	AL121753	1.30
autocrine motility factor receptor	NM_001144	1.30
tenascin XB	BE044614	1.30
adenosine deaminase	Z97053	1.30
nuclear factor of activated T-cells, cytoplasmic, calcineurin-dependent 3	NM_004555	1.30
fracture callus 1 homolog (rat)	NM_012192	1.30
KIAA0514 gene product	NM_014696	1.30
putative ATPase	NM_022089	1.30
Ras suppressor protein 1	NM_012425	1.30
translation factor sui1 homolog	NM_005875	1.30
lectin, galactoside-binding, soluble, 3 binding protein	NM_005567	1.30
hypothetical protein	AL136892	1.30
CDC10 cell division cycle 10 homolog (S. cerevisiae)	BE551361	1.30
Consensus includes gb:AF070536.1 /DEF=Homo sapiens clone 24566 mRNA sequence. /FEA=mRNA /DB_XREF=gi:3387893 /UG=Hs.133342 Homo sapiens clone 24566 mRNA sequence	AF070536	1.30
nuclear receptor subfamily 2, group F, member 1	BC004154	1.30
carcinoembryonic antigen-related cell adhesion molecule 7	L31792	1.30
potential tumor suppressor	NM_013437	1.30
diacylglycerol kinase, alpha (80kD)	NM_001345	1.30
DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 31	NM_022779	1.30
elastase 1, pancreatic	NM_001971	1.30
small inducible cytokine subfamily B (Cys-X-Cys), member 14 (BRAK)	NM_004887	1.30
basic transcription factor 3, like 1	NM_001208	1.30
mitogen-activated protein kinase kinase 6	U39657	1.30
hypothetical protein FLJ23053	NM_022907	1.30
hypothetical protein	NM_019018	1.30
Consensus includes gb:AK024944.1 /DEF=Homo sapiens cDNA: FLJ21291 fis, clone COL01963. /FEA=mRNA /DB_XREF=gi:10437366 /UG=Hs.287657 Homo sapiens cDNA: FLJ21291 fis, cl	AK024944	1.30
plexin B3	NM_005393	1.30
spinocerebellar ataxia 7 (olivopontocerebellar atrophy with retinal degeneration)	AF032105	1.30
adducin 1 (alpha)	AL556041	1.30
ectonucleotide pyrophosphatase/phosphodiesterase 1	NM_006208	1.30
hypothetical protein FLJ00001	BC000122	1.30
huntingtin-associated protein interacting protein (duo)	NM_003947	1.30
splicing factor, arginine/serine-rich 3	AF107405	1.30
KIAA0673 protein	AL117405	1.30
6 protein-coupled receptor 49	AL524520	1.30
dopa decarboxylase (aromatic L-amino acid decarboxylase)	NM_000790	1.30
PRO1914 protein	NM_014106	1.30
Consensus includes gb:AK026238.1 /DEF=Homo sapiens cDNA: FLJ22585 fis, clone HSI02678. /FEA=mRNA /DB_XREF=gi:10439027 /UG=Hs.287701 Homo sapiens cDNA: FLJ22585 fis, cl	AK026238	1.30
hypothetical protein PRO0386	NM_018562	1.30
endothelin converting enzyme 1	BF969352	1.30
hypothetical protein PRO3073	NM_018549	1.30

opioid receptor, mu 1	L29301	1.30
t-complex 10 (a murine tcp homolog)	AL021331	1.30
putative homeodomain transcription factor 1	NM_006608	1.30
C-reactive protein, pentraxin-related	X56692	1.30
hydroxyprostaglandin dehydrogenase 15-(NAD)	U63296	1.30
peptide transporter 3	NM_016582	1.30
Consensus includes gb:AL137284.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434D1516 (from clone DKFZp434D1516). /FEA=mRNA /DB_XREF=gi:6807739 /UG=Hs.274572 Homo sapiens	AL137284	1.30
interleukin-1 homolog 1	NM_019618	1.30
sel-1 suppressor of lin-12-like (C. elegans)	AI927770	1.30
capping protein (actin filament) muscle Z-line, alpha 1	NM_006135	1.30
KIAA0964 protein	AW134976	1.30
hypothetical protein FLJ10488	NM_018110	1.30
suppression of tumorigenicity 5	NM_005418	1.30
hypothetical protein FLJ14050	NM_024869	1.30
casein, alpha	NM_001890	1.30
proteasome (prosome, macropain) 26S subunit, ATPase, 4	AC007842	1.30
histidine decarboxylase	NM_002112	1.30
zinc finger protein 81 (HFZ20)	X68011	1.30
Yes-associated protein 1, 65 kDa	AI745185	1.30
RAB11B, member RAS oncogene family	X79780	1.30
STAT induced STAT inhibitor-4	NM_004232	1.30
solute carrier family 9 (sodium/hydrogen exchanger), isoform 5	NM_004594	1.30
prostaglandin I2 (prostacyclin) synthase	D38145	1.30
zinc finger protein 154 (pHZ-92)	U20648	1.30
bradykinin receptor B2	NM_000623	1.30
Consensus includes gb:BG236351 /FEA=EST /DB_XREF=gi:12750198 /DB_XREF=est:naf26f12.x1 /CLONE=IMAGE:4142350 /UG=Hs.148161 ESTs	BG236351	1.30
annexin A4	BC000182	1.30
TAF2 RNA polymerase II, TATA box binding protein (TBP)-associated factor, 150 kD	AK001618	1.30
cellular retinoic acid binding protein 1	NM_004378	1.30
topoisomerase (DNA) I	J03250	1.30
sin3-associated polypeptide, 30kD	NM_003864	1.30
Consensus includes gb:AL117652.1 /DEF=Homo sapiens mRNA; cDNA DKFZp586B1324 (from clone DKFZp586B1324). /FEA=mRNA /DB_XREF=gi:5912244 /UG=Hs.278334 Homo sapiens	AL117652	1.30
caspase recruitment domain family, member 15	NM_022162	1.30
Consensus includes gb:AL137495.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434C2331 (from clone DKFZp434C2331). /FEA=mRNA /DB_XREF=gi:6808119 /UG=Hs.306468 Homo sapiens	AL137495	1.30
semaphorin sem2	NM_020163	1.30
activity-regulated cytoskeleton-associated protein	AF193421	1.30
a disintegrin-like and metalloprotease (reprolysin type) with thrombospondin type 1 motif, 2	NM_014244	1.30
fibroblast growth factor 18	BC006245	1.30
calbindin 1, (28kD)	AW014927	1.30
solute carrier family 7 (cationic amino acid transporter, y+ system), member 2	NM_003046	1.30
hypothetical protein FLJ20051	NM_019087	1.30
ACO for serine protease homologue	AF243527	1.30
neuropeptide Y receptor Y2	U36269	1.30
Consensus includes gb:AW574818 /FEA=EST /DB_XREF=gi:7246357 /DB_XREF=est:UI-HF-BK0-abi-e-06-0-UI.s1 /CLONE=IMAGE:3056362 /UG=Hs.32374 ESTs, Weakly similar to KIAA1528	AW574818	1.30
argininosuccinate synthetase	NM_000050	1.30
solute carrier family 35 (UDP-galactose transporter), member 2	NM_005660	1.30
unc-119 homolog (C. elegans)	NM_005148	1.30
transmembrane protease, serine 3	NM_024022	1.30
signal transducer and activator of transcription 5A	NM_003152	1.30
chloride channel 2	AI141670	1.30
coagulation factor XII (Hageman factor)	BF530348	1.30
hypothetical protein FLJ10698	AL050219	1.30
KIAA1199 protein	AB033025	1.30
PTH-responsive osteosarcoma B1 protein	U87408	1.30
heat shock 27kD protein family, member 7 (cardiovascular)	NM_014424	1.30
dopamine receptor D3	NM_000796	1.30
hypothetical protein DKFZp434A1721	NM_017609	1.30
putative nuclear protein	AA046650	1.30
calcium binding protein 5	NM_019855	1.30
heat shock protein (hsp110 family)	NM_014278	1.30
short stature homeobox	NM_000451	1.30
eukaryotic translation initiation factor 3, subunit 5 (epsilon, 47kD)	AI001896	1.30
dynein, axonemal, heavy polypeptide 17	AL122077	1.30

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synovial sarcoma translocation, chromosome 18	AF257501	1.30
matrix metalloproteinase 9 (gelatinase B, 92kD gelatinase, 92kD type IV collagenase)	NM_004994	1.30
par-6 partitioning defective 6 homolog beta (C. elegans)	AB044555	1.30
protein phosphatase 1B (formerly 2C), magnesium-dependent, beta isoform	AJ271832	1.30
nuclear receptor subfamily 4, group A, member 2	AJ935096	1.30
reversion-inducing-cysteine-rich protein with kazal motifs	NM_021111	1.30
Consensus includes gb:BF542055 /FEA=EST /DB_XREF=gi:11629436 /DB_XREF=est:602068320F1 /CLONE=IMAGE:4067357 /UG=Hs.190785 ESTs	BF542055	1.30
transcription factor 3 (E2A immunoglobulin enhancer binding factors E12/E47)	AI871234	1.30
Consensus includes gb:AK025862.1 /DEF=Homo sapiens cDNA: FLJ22209 fis, clone HRC01496. /FEA=mRNA /DB_XREF=gi:10438505 /UG=Hs.288528 Homo sapiens cDNA: FLJ22209 fis, cl	AK025862	1.30
valosin-containing protein	W60953	1.30
ring finger protein 1	AL031228	1.30
glucokinase (hexokinase 4) regulatory protein	NM_001486	1.30
M27830 Human 28S ribosomal RNA gene, complete cds (_5, _M, _3 represent transcript regions 5 prime, Middle, and 3 prime respectively)	M27830	1.30
CGI-116 protein	NM_016053	1.30
protease inhibitor 3, skin-derived (SKALP)	NM_002638	1.30
annexin A13	NM_004306	1.30
KIAA0843 protein	NM_014945	1.30
adenylate cyclase activating polypeptide 1 (pituitary)	NM_001117	1.30
cyclin F	U17105	1.30
EphA3	AF213460	1.30
fetuin B	AB017551	1.30
retinal homeobox protein	NM_013435	1.30
keratin, hair, acidic, 7	NM_003770	1.30
Consensus includes gb:AU158566 /FEA=EST /DB_XREF=gi:11020087 /DB_XREF=est:AU158566 /CLONE=PLACE3000322 /UG=Hs.287635 Homo sapiens cDNA FLJ14312 fis, clone PLACE3(	AU158566	1.30
Consensus includes gb:BE672818 /FEA=EST /DB_XREF=gi:10033359 /DB_XREF=est:7d24c01.x1 /CLONE=IMAGE:3248160 /UG=Hs.297681 serine (or cysteine) proteinase inhibitor, clade	BE672818	1.30
major histocompatibility complex, class I-like sequence	NM_001531	1.30
transcription factor 20 (AR1)	AB006630	1.30
geminin	NM_015895	1.30
myosin VIIA (Usher syndrome 1B (autosomal recessive, severe))	U39226	1.30
Consensus includes gb:AW242701 /FEA=EST /DB_XREF=gi:6576546 /DB_XREF=est:xm90a08.x1 /CLONE=IMAGE:2691446 /UG=Hs.256398 Homo sapiens mRNA; cDNA DKFZp434E0528 (fi	AW242701	1.30
CD28 antigen (Tp44)	AF222343	1.30
Homer, neuronal immediate early gene, 2	Y19026	1.30
calcium/calmodulin-dependent protein kinase (CaM kinase) II alpha	AW131813	1.30
Consensus includes gb:AK024381.1 /DEF=Homo sapiens cDNA FLJ14319 fis, clone PLACE3000406. /FEA=mRNA /DB_XREF=gi:10436753 /UG=Hs.287636 Homo sapiens cDNA FLJ14319 fi:	AK024381	1.30
TAF9-like RNA polymerase II, TATA box binding protein (TBP)-associated factor, 31 kD	AF220509	1.30
hypothetical protein MGC11352	NM_030927	1.30
Consensus includes gb:AI143879 /FEA=EST /DB_XREF=gi:3665688 /DB_XREF=est:qe04f11.x1 /CLONE=IMAGE:1738029 /UG=Hs.128291 phosphodiesterase 10A /FL=gb:AF127479.1 gb:/	AI143879	1.30
connexin46.6	BF061637	1.30
histamine receptor H1	D28481	1.30
hypothetical protein FLJ10040	NM_017977	1.30
Consensus includes gb:AL049313.1 /DEF=Homo sapiens mRNA; cDNA DKFZp564B076 (from clone DKFZp564B076). /FEA=mRNA /DB_XREF=gi:4500086 /UG=Hs.21103 Homo sapiens mRI	AL049313	1.30
doublecortin and CaM kinase-like 1	NM_004734	1.30
Consensus includes gb:BG149837 /FEA=EST /DB_XREF=gi:12661867 /DB_XREF=est:nae01b12.x1 /CLONE=IMAGE:3434062 /UG=Hs.33540 ESTs, Weakly similar to dJ309K20.4 H.sapiens	BG149837	1.30
RNA guanylyltransferase and 5'-phosphatase	AB012143	1.30
hypothetical protein FLJ12895	NM_023926	1.30
monoamine oxidase A	NM_000240	1.30
hydroxyacid oxidase 2 (long chain)	NM_016527	1.30
adenylate cyclase 1 (brain)	L05500	1.30
myosin X	AJ561354	1.30
KIAA1237 protein	AA121502	1.30
Consensus includes gb:AK022309.1 /DEF=Homo sapiens cDNA FLJ12247 fis, clone MAMMA1001397. /FEA=mRNA /DB_XREF=gi:10433678 /UG=Hs.287506 Homo sapiens cDNA FLJ12247 f	AK022309	1.30
hypothetical protein FLJ12584	NM_025139	1.30
mitochondrial ribosomal protein 63	BF195165	1.30
Niemann-Pick disease, type C1	NM_000271	1.30
Consensus includes gb:AK026449.1 /DEF=Homo sapiens cDNA: FLJ22796 fis, clone KAIA2544. /FEA=mRNA /DB_XREF=gi:10439318 /UG=Hs.306850 Homo sapiens cDNA: FLJ22796 fis, clc	AK026449	1.30
ubiquitously transcribed tetratricopeptide repeat gene, X chromosome	AI140752	1.30
paired mesoderm homeo box 1	NM_006902	1.30
STAT induced STAT inhibitor-2	NM_003877	1.30
peptidylarginine deiminase type III	NM_016233	1.30
secretogranin II (chromogranin C)	NM_003469	1.30
zinc finger protein 262	NM_005095	1.29
biliverdin reductase A	BC005902	1.29
keratin 18	NM_000224	1.29

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ribosomal protein S6 kinase, 52kD, polypeptide 1	NM_012424	1.29
KIAA1105 protein	N38985	1.29
cytoplasmic linker 2	BC006259	1.29
calpain 2, (m/II) large subunit	AK023851	1.29
chromosome 5 open reading frame 7	BC001202	1.29
Consensus includes gb:AK025188.1 /DEF=Homo sapiens cDNA: FLJ21535 fis, clone COL06131. /FEA=mRNA /DB_XREF=gi:10437651 /UG=Hs.306781 Homo sapiens cDNA: FLJ21535 fis, cl	AK025188	1.29
gb:BC005914.1 /DEF=Homo sapiens, clone MGC:14510, mRNA, complete cds. /FEA=mRNA /PROD=Unknown (protein for MGC:14510) /DB_XREF=gi:13543511 /FL=gb:BC005914.1	BC005914	1.29
FT005 protein	NM_014054	1.29
caspase 10, apoptosis-related cysteine protease	AF111344	1.29
Consensus includes gb:AL049369.1 /DEF=Homo sapiens mRNA; cDNA DKFZp586D0518 (from clone DKFZp586D0518). /FEA=mRNA /DB_XREF=gi:4500160 /UG=Hs.250724 Homo sapiens	AL049369	1.29
hypothetical protein FKS628	NM_030929	1.29
hypothetical protein FLJ23403	NM_022068	1.29
KIAA0887 protein	AB020694	1.29
gamma-aminobutyric acid (GABA) A receptor, beta 2	NM_021911	1.29
fibulin 2	NM_001998	1.29
striatin, calmodulin binding protein	NM_003162	1.29
insulin-like growth factor 1 receptor	H05812	1.29
TAF5 RNA polymerase II, TATA box binding protein (TBP)-associated factor, 100 kD	AW138827	1.29
HCF-binding transcription factor Zhangfei	AI762013	1.29
hypothetical protein FLJ20300	NM_017753	1.29
phosphoinositol 3-phosphate-binding protein-2	NM_019012	1.29
KIAA0352 gene product	NM_014830	1.29
microseminoprotein, beta-	U22178	1.29
interferon regulatory factor 1	NM_002198	1.29
Consensus includes gb:AL137673.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434H0872 (from clone DKFZp434H0872). /FEA=mRNA /DB_XREF=gi:6807841 /UG=Hs.306454 Homo sapiens	AL137673	1.29
ELL gene (11-19 lysine-rich leukemia gene)	AL521391	1.29
oral-facial-digital syndrome 1	NM_003611	1.29
polyadenylate binding protein-interacting protein 1	AW613203	1.29
hypothetical protein FLJ13241	NM_025088	1.29
lipocalin-interacting membrane receptor	NM_018113	1.29
Consensus includes gb:AA127885 /FEA=EST /DB_XREF=gi:1687192 /DB_XREF=est:zl13b04.s1 /CLONE=IMAGE:501775 /UG=Hs.13350 Homo sapiens mRNA; cDNA DKFZp586D0918 (from	AA127885	1.29
KIAA0484 protein	AA732995	1.29
polyadenylate binding protein-interacting protein 1	BC005295	1.29
SCAN domain-containing 2	AF244812	1.29
muscle disease-related protein	NM_016298	1.29
GRB2-related adaptor protein 2	NM_004810	1.29
cytochrome c oxidase subunit IV isoform 1	AW337510	1.29
interleukin 18 receptor 1	NM_003855	1.29
thyroid peroxidase	M17755	1.29
hypothetical protein PRO2214	NM_018517	1.29
KIAA0657 protein	AI915513	1.29
deiodinase, iodothyronine, type II	AB041844	1.29
RAB3B, member RAS oncogene family	BC005035	1.29
splicing factor proline/glutamine rich (polypyrimidine tract binding protein associated)	BG035151	1.29
KIAA0470 gene product	NM_014812	1.29
solute carrier family 6 (neurotransmitter transporter, GABA), member 11	NM_014229	1.29
fibroblast growth factor receptor 3 (achondroplasia, thanatophoric dwarfism)	NM_000142	1.29
Consensus includes gb:AI718937 /FEA=EST /DB_XREF=gi:5036193 /DB_XREF=est:as50b04.x1 /CLONE=IMAGE:2320591 /UG=Hs.109438 Homo sapiens clone 24775 mRNA sequence	AI718937	1.29
stromal cell derived factor receptor 1	NM_017455	1.29
zinc finger protein 36, C3H type-like 1	AI344045	1.29
alcohol dehydrogenase 7 (class IV), mu or sigma polypeptide	U07821	1.29
myosin light chain 2a	NM_021223	1.29
Consensus includes gb:AL035406 /DEF=Human DNA sequence from clone RP1-233K16 on chromosome 1p36.21-36.33 Contains the gene KIAA0444, a putative chromodomain helicase DNA	AL035406	1.29
6 protein-coupled receptor 51	AF095723	1.29
ATP synthase, H+ transporting, mitochondrial F1 complex, gamma polypeptide 1	BG232034	1.29
DIPB protein	AA176780	1.29
S-adenosylmethionine decarboxylase 1	NM_001634	1.29
T-box 3 (ulnar mammary syndrome)	NM_016569	1.29
receptor-associated protein of the synapse, 43kD	BC004196	1.29
collagen, type IV, alpha 6	BC005305	1.29
gamma-glutamyltransferase 1	NM_013430	1.29
activator of S phase kinase	NM_006716	1.29
Consensus includes gb:AL109707.1 /DEF=Homo sapiens mRNA full length insert cDNA clone EUROIMAGE 295344. /FEA=mRNA /DB_XREF=gi:5689824 /UG=Hs.94327 Homo sapiens mRNA	AL109707	1.29

a disintegrin and metalloproteinase domain 28	NM_021778	1.29
KIAA1067 protein	AI817079	1.29
carcinoembryonic antigen-related cell adhesion molecule 3	NM_001815	1.29
SGC32445 protein	NM_016606	1.29
KIAA0731 protein	BC001460	1.29
discs, large (Drosophila) homolog-associated protein 1	NM_004746	1.29
dynein, cytoplasmic, light intermediate polypeptide 2	AI640861	1.29
proline-rich protein HaeIII subfamily 1	NM_006250	1.29
Cluster Incl. N53536:yz26h01.s1 Homo sapiens cDNA, 3 end /clone=IMAGE-284209 /clone_end=3 /gb=N53536 /gi=1194702 /ug=Hs.33540 /len=543	N53536	1.29
Consensus includes gb:U52428 /DEF=Human fatty acid synthase gene, partial cds /FEA=CDS_2 /DB_XREF=gi:1374810 /UG=Hs.83190 fatty acid synthase	U52428	1.29
MSTP032 protein	NM_025226	1.29
egf-like module containing, mucin-like, hormone receptor-like sequence 2	NM_013447	1.29
KIAA0408 gene product	NM_014702	1.29
protocadherin gamma subfamily B, 5	AF152521	1.29
TAF10 RNA polymerase II, TATA box binding protein (TBP)-associated factor, 30 kD	NM_006284	1.29
Rhesus blood group, B glycoprotein	NM_020407	1.29
KIAA0068 protein	AK026548	1.29
cell division cycle 2, G1 to S and G2 to M	AL524035	1.29
reticulon 4	AF333336	1.29
hypothetical protein FLJ20950	NM_024952	1.29
pyruvate dehydrogenase kinase, isoenzyme 2	AI870615	1.29
collagen, type XIII, alpha 1	M33653	1.29
MCF.2 cell line derived transforming sequence	X13230	1.29
HMT1 hnRNP methyltransferase-like 3 (S. cerevisiae)	NM_019854	1.29
Consensus includes gb:AL137706.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434A2322 (from clone DKFZp434A2322). /FEA=mRNA /PROD=hypothetical protein /DB_XREF=gi:6808070 /UG	AL137706	1.29
zinc finger protein 79 (pT7)	AA284829	1.29
Kruppel-like factor 1 (erythroid)	U65404	1.29
ubiquitin-conjugating enzyme E2A (RAD6 homolog)	AI126625	1.29
Wiskott-Aldrich syndrome-like	NM_003941	1.29
glutamate receptor, ionotropic, kainate 1	U16125	1.29
glucose-6-phosphate dehydrogenase	X55448	1.29
myeloproliferative leukemia virus oncogene	M90103	1.29
dystrobrevin, alpha	NM_001390	1.29
regulator of G-protein signalling 12	AI824113	1.29
Consensus includes gb:AI478300 /FEA=EST /DB_XREF=gi:4371526 /DB_XREF=est:tm39e01.x1 /CLONE=IMAGE:2160504 /UG=Hs.192789 ESTs, Weakly similar to ALU6_HUMAN ALU SUBFA	AI478300	1.29
SH2 domain-containing phosphatase anchor protein 1	NM_030764	1.29
Kelch motif containing protein	BC005253	1.29
GA binding protein transcription factor, beta subunit 1 (53kD)	NM_005254	1.29
KIAA0427 gene product	NM_014772	1.29
T cell receptor alpha locus	BF976764	1.29
peroxiredoxin 3	NM_006793	1.29
phosphoinositide-specific phospholipase C-beta 1	AA393484	1.29
phosphoinositol 3-phosphate-binding protein-2	NM_014935	1.29
Consensus includes gb:BE541042 /FEA=EST /DB_XREF=gi:9769786 /DB_XREF=est:601064390F1 /CLONE=IMAGE:3450599 /UG=Hs.23240 Homo sapiens cDNA FLJ13496 fis, clone PLACE	BE541042	1.29
acyl-Coenzyme A dehydrogenase, long chain	AI367275	1.29
pan-hematopoietic expression	NM_005705	1.29
G protein-coupled receptor 39	AL567376	1.29
KIAA1193 protein	AW003889	1.29
natriuretic peptide receptor A/guanylate cyclase A (atrionatriuretic peptide receptor A)	NM_000906	1.29
metastasis-associated 1-like 1	AB012922	1.29
heterogeneous nuclear ribonucleoprotein D (AU-rich element RNA binding protein 1, 37kD)	NM_002138	1.29
C-terminal binding protein 2	AF22271.1	1.29
hypothetical protein FLJ11160	NM_018344	1.29
Consensus includes gb:AK026106.1 /DEF=Homo sapiens cDNA: FLJ22453 fis, clone HRC09679, highly similar to AF059516 Homo sapiens tolloid-like 2 protein (TLL2) mRNA. /FEA=mRNA /	AK026106	1.29
proteoglycan 1, secretory granule	J03223	1.29
TAF1 RNA polymerase II, TATA box binding protein (TBP)-associated factor, 250 kD	M73444	1.29
fatty-acid-Coenzyme A ligase, very long-chain 1	NM_003645	1.29
gamma-glutamyltransferase 1	L20490	1.29
Consensus includes gb:AU146791 /FEA=EST /DB_XREF=gi:11008312 /DB_XREF=est:AU146791 /CLONE=HEMBB1001531 /UG=Hs.287474 Homo sapiens cDNA FLJ12000 fis, clone HEMBB	AU146791	1.29
guanine nucleotide binding protein (G protein), gamma transducing activity polypeptide 1	NM_021955	1.29
adenosine monophosphate deaminase 2 (isoform L)	AI916249	1.29
spinocerebellar ataxia 1 (olivopontocerebellar ataxia 1, autosomal dominant, ataxin 1)	AW235612	1.29
cofactor required for Sp1 transcriptional activation, subunit 2 (150kD)	BE567032	1.29

hypothetical protein FLJ21616	NM_024567	1.29
Consensus includes gb:AI207120 /FEA=EST /DB_XREF=gi:3765792 /DB_XREF=est:qf59e09.x1 /CLONE=IMAGE:1754344 /UG=Hs.248129 glutathione peroxidase 5 (epididymal androgen-r	AI207120	1.29
potassium intermediate/small conductance calcium-activated channel, subfamily N, member 2	NM_021614	1.29
cardiotrophin-like cytokine; neurotrophin-1/B-cell stimulating factor-3	NM_013246	1.29
FN5 protein	NM_020179	1.29
potassium large conductance calcium-activated channel, subfamily M, beta member 3-like	NM_014406	1.29
surfactant protein B	L11573	1.29
aspartate beta-hydroxylase	NM_020164	1.29
Consensus includes gb:L37198.1 /DEF=Homo sapiens (clone B3B3E13) Huntingtons disease candidate region mRNA fragment. /FEA=mRNA /DB_XREF=gi:600519 /UG=Hs.233617 Homo :	L37198	1.29
KIAA1698 protein	AI869988	1.29
H3 histone family, member F	NM_003533	1.29
KIAA0429 gene product	AI027678	1.29
prostaglandin E synthase	AF010316	1.29
fibrinogen, B beta polypeptide	BG545288	1.29
insulin receptor substrate 3-like	H07885	1.29
kallikrein 8 (neuropsin/ovasin)	NM_007196	1.29
leukocyte immunoglobulin-like receptor, subfamily B (with TM and ITIM domains), member 5	NM_006840	1.29
v-ros UR2 sarcoma virus oncogene homolog 1 (avian)	NM_002944	1.29
small inducible cytokine subfamily A (Cys-Cys), member 13	NM_005408	1.29
v-rel reticuloendotheliosis viral oncogene homolog A, nuclear factor of kappa light polypeptide gene enhancer in B-cells 3, p65 (avian)	M62399	1.29
general transcription factor IIIC, polypeptide 2 (beta subunit, 110kD)	NM_001521	1.29
type I transmembrane protein Fn14	NM_016639	1.29
CGI-105 protein	AA723370	1.29
angiotensin receptor 2	U16957	1.29
hypothetical protein	NM_015677	1.29
melatonin receptor 1A	NM_005958	1.29
keratin 4	BF588953	1.29
testes-epecific heterogenous nuclear ribonucleoprotein G-T	AF069682	1.29
DKFZP564N1363 protein	AL121895	1.29
tumor necrosis factor receptor superfamily, member 4	AI277151	1.29
ventral anterior homeobox 2	NM_012476	1.29
protein phosphatase 2, regulatory subunit B (B56), beta isoform	NM_006244	1.29
prostaglandin E receptor 3 (subtype EP3)	L27489	1.29
oculocerebrorenal syndrome of Lowe	M88162	1.29
MADS box transcription enhancer factor 2, polypeptide B (myocyte enhancer factor 2B)	BC004449	1.29
calcitonin receptor-like	NM_005795	1.29
synovial sarcoma, X breakpoint 5	NM_021015	1.29
KIAA1025 protein	AK023837	1.29
G protein-coupled receptor 32	NM_001506	1.29
hypothetical protein FLJ23495	NM_024800	1.29
homeo box B8	NM_024016	1.29
KIAA1277 protein	AI131051	1.29
reticulon 1	BC000314	1.29
neuropilin 2	AF022859	1.29
nuclear receptor subfamily 1, group I, member 2	AF061056	1.29
Consensus includes gb:AL137624.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434M1812 (from clone DKFZp434M1812). /FEA=mRNA /DB_XREF=gi:6808420 /UG=Hs.306476 Homo sapiens	AL137624	1.29
enolase 3, (beta, muscle)	NM_001976	1.29
myelin transcription factor 2	NM_003871	1.29
lymphocyte antigen 95 (activating NK-receptor; NK-p44)	NM_004828	1.29
centaurin, beta 1	NM_014716	1.29
transcription factor (SMIF gene)	NM_018403	1.29
KIAA0914 gene product	AW450403	1.29
EGF-containing fibulin-like extracellular matrix protein 1	NM_004105	1.29
KIAA0752 protein	AW514267	1.29
cyclin D1 (PRAD1: parathyroid adenomatosis 1)	Z23022	1.29
Fanconi anemia, complementation group G	AC004472	1.29
STE20-like kinase	AF181985	1.29
cyclin D1 (PRAD1: parathyroid adenomatosis 1)	M73554	1.29
chondroitin sulfate proteoglycan 3 (neurocan)	NM_004386	1.29
myosin, heavy polypeptide 2, skeletal muscle, adult	NM_017534	1.29
Consensus includes gb:AF195953 /DEF=Homo sapiens membrane-bound aminopeptidase P (XNPEP2) gene, complete cds /FEA=mRNA /DB_XREF=gi:11066156 /UG=Hs.57922 X-prolyl am	AF195953	1.29
KIAA0326 protein	AI924293	1.29
Consensus includes gb:AA757368 /FEA=EST /DB_XREF=gi:2805231 /DB_XREF=est:ah98a10.s1 /CLONE=IMAGE:1327098 /UG=Hs.7057 Homo sapiens Cri-du-chat region mRNA, clone NIBI	AA757368	1.29

laminin, beta 2 (laminin S)	X79683	1.29
optic atrophy 1 (autosomal dominant)	AA209332	1.29
hepcidin antimicrobial peptide	NM_021175	1.29
sirtuin silent mating type information regulation 2 homolog 2 (S. cerevisiae)	NM_012237	1.29
chromosome 5 open reading frame 3	NM_018691	1.29
hypothetical protein FLJ10853	NM_018246	1.29
Consensus includes gb:AI523613 /FEA=EST /DB_XREF=gi:4437748 /DB_XREF=est:tg95a07.x1 /CLONE=IMAGE:2116500 /UG=Hs.293495 ESTs, Weakly similar to ALU1_HUMAN ALU SUBFA	AI523613	1.29
proteinzr0008	NM_013395	1.29
hypothetical protein FLJ11017	NM_018302	1.29
hypothetical protein FLJ14281	NM_024920	1.29
poly(rC) binding protein 3	NM_020528	1.29
Consensus includes gb:AL136527 /DEF=Human DNA sequence from clone RP11-215B13 on chromosome 13 Contains the AKAP11 (A kinase (PRKA) anchor protein 11) gene, part of a gene	AL136527	1.29
6-phosphofructo-2-kinase/fructose-2,6-biphosphatase 2	AB044805	1.29
synaptonemal complex protein 1	NM_003176	1.29
methylthioadenosine phosphorylase	AF109294	1.29
interferon regulatory factor 4	D78261	1.29
cancer/testis antigen	AI275978	1.29
hypothetical protein DKFZp564F013	AL050389	1.29
homeodomain interacting protein kinase 3	AF305239	1.28
Consensus includes gb:AK024355.1 /DEF=Homo sapiens cDNA FLJ14293 fis, clone PLACE1007866. /FEA=mRNA /DB_XREF=gi:10436719 /UG=Hs.141009 Homo sapiens cDNA FLJ14293 fi	AK024355	1.28
KIAA0329 gene product	AB002295	1.28
ligand of neuronal nitric oxide synthase with carboxyl-terminal PDZ domain	AF037070	1.28
NAG-5 protein	NM_016446	1.28
importin 13	NM_014652	1.28
olfactory receptor, family 1, subfamily F, member 2	AJ003145	1.28
ret finger protein-like 1	NM_021026	1.28
solute carrier family 21 (organic anion transporter), member 3	AF085224	1.28
glia maturation factor, beta	NM_004124	1.28
gb:U50073.1 /DEF=Human T cell receptor beta chain precursor (Vbeta 12.3) mRNA, complete cds. /FEA=mRNA /GEN=Vbeta 12.3 /PROD=T cell receptor beta chain precursor /DB_XREF=g	U50073	1.28
estrogen receptor 1	AF258450	1.28
amnionless protein	NM_030943	1.28
uridine monophosphate synthetase (orotate phosphoribosyl transferase and orotidine-5'-decarboxylase)	AL080099	1.28
KIAA0579 protein	AL117532	1.28
integrin, alpha X (antigen CD11C (p150), alpha polypeptide)	M81695	1.28
cytochrome P450, subfamily IIIA (niphedipine oxidase), polypeptide 3	J04449	1.28
TBP-interacting protein	AB014567	1.28
carbohydrate (chondroitin 6/keratan) sulfotransferase 3	AB017915	1.28
inter-alpha (globulin) inhibitor, H1 polypeptide	AF116713	1.28
protein phosphatase 2, regulatory subunit B (B56), beta isoform	L42374	1.28
olfactory receptor, family 1, subfamily A, member 2	NM_012352	1.28
mesenchyme homeo box 2 (growth arrest-specific homeo box)	NM_005924	1.28
ATP-binding cassette, sub-family C (CFTR/MRP), member 2	NM_000392	1.28
double C2-like domains, alpha	NM_003586	1.28
zinc finger protein 93 homolog (mouse)	NM_004234	1.28
alkaline phosphatase, placental-like 2	X07247	1.28
apolipoprotein D	NM_001647	1.28
cofactor required for Sp1 transcriptional activation, subunit 2 (150kD)	NM_004229	1.28
f-box and leucine-rich repeat protein 2	AF176518	1.28
homeo box C11	NM_014212	1.28
Consensus includes gb:AL356954 /DEF=Human DNA sequence from clone RP11-520F24 on chromosome 13 Contains ESTs, STSs and GSSs. Contains an HNRPA1 (heterogenous nuclear ribo	AL356954	1.28
HERV-H LTR-associating 1	NM_005712	1.28
KIAA0444 protein	AL110281	1.28
Consensus includes gb:AI655015 /FEA=EST /DB_XREF=gi:4738994 /DB_XREF=est:wb66a10.x1 /CLONE=IMAGE:2310618 /UG=Hs.3843 Homo sapiens mRNA; cDNA DKFZp586F2224 (from	AI655015	1.28
protocadherin alpha 2	AF152480	1.28
nuclear receptor subfamily 1, group I, member 2	NM_003889	1.28
HIC1-related gene on chromosome 22	AL043112	1.28
hypothetical protein FLJ11850	NM_022741	1.28
gb:U41742.1 /DEF=Human nucleophosmin-retinoic acid receptor alpha fusion protein NPM-RAR long form mRNA, complete cds. /FEA=mRNA /PROD=nucleophosmin-retinoic acid receptor a	U41742	1.28
hypothetical protein	NM_015372	1.28
Consensus includes gb:BG536504 /FEA=EST /DB_XREF=gi:13528050 /DB_XREF=est:602564909F1 /CLONE=IMAGE:4689314 /UG=Hs.274518 Homo sapiens mRNA; cDNA DKFZp586O202	BG536504	1.28
cytochrome b5 outer mitochondrial membrane precursor	AF130113	1.28
CD80 antigen (CD28 antigen ligand 1, B7-1 antigen)	NM_005191	1.28
hypothetical protein FLJ21140	NM_024776	1.28

v-fos FBJ murine osteosarcoma viral oncogene homolog	BC004490	1.28
Consensus includes gb:S83374.1 /DEF=glutamate transporter II variant BHB6T IIB (5 region) human, brain and spinal cord, mRNA Partial Mutant, 129 nt. /FEA=mRNA /DB_XREF=gi:191162	S83374	1.28
signal-regulatory protein beta 2	NM_018556	1.28
Consensus includes gb:AL583528 /FEA=EST /DB_XREF=gi:12952578 /DB_XREF=est:AL583528 /CLONE=CS0DD004YH21 (5 prime) /UG=Hs.42915 ARP2 (actin-related protein 2, yeast) hc	AL583528	1.28
signal transducer and activator of transcription 1, 91kD	M97935	1.28
SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily f, member 1	NM_006015	1.28
Consensus includes gb:AF090883.1 /DEF=Homo sapiens clone HQ0021. /FEA=mRNA /DB_XREF=gi:6690149 /UG=Hs.306560 Homo sapiens clone HQ0021	AF090883	1.28
integrin, alpha 10	AF112345	1.28
solute carrier family 9 (sodium/hydrogen exchanger), isoform 1 (antiporter, Na+/H+, amiloride sensitive)	M81768	1.28
chromosome 11 open reading frame 21	NM_014144	1.28
LIM homeobox protein 3	AF156888	1.28
cyclin-dependent kinase-like 3	NM_016508	1.28
death effector domain-containing	NM_004216	1.28
collagen, type V, alpha 1	AI130969	1.28
eyes absent homolog 1 (Drosophila)	AJ000098	1.28
KIAA0690 protein	AK021460	1.28
hypothetical protein FLJ10647	NM_018166	1.28
signal transducer and activator of transcription 1, 91kD	M97935	1.28
transcription factor 3 (E2A immunoglobulin enhancer binding factors E12/E47)	M31222	1.28
protein phosphatase 1, regulatory (inhibitor) subunit 13B	AK023188	1.28
ras-related C3 botulinum toxin substrate 3 (rho family, small GTP binding protein Rac3)	NM_005052	1.28
phosphoinositide-3-kinase, class 2, beta polypeptide	NM_002646	1.28
protocadherin gamma subfamily C, 3	AF152524	1.28
interleukin 15 receptor, alpha	NM_002189	1.28
mitogen-activated protein kinase kinase kinase kinase 3	NM_003618	1.28
nucleoporin 50kD	AF267865	1.28
caspase 1, apoptosis-related cysteine protease (interleukin 1, beta, convertase)	U13700	1.28
aldehyde dehydrogenase 9 family, member A1	NM_000696	1.28
BRF2, subunit of RNA polymerase III transcription initiation factor, BRF1-like	AF298153	1.28
transmembrane gamma-carboxyglutamic acid protein 3	NM_024082	1.28
cholinergic receptor, nicotinic, beta polypeptide 3	NM_000749	1.28
hepatocellular carcinoma-associated antigen 127	NM_018684	1.28
paraneoplastic cancer-testis-brain antigen	NM_013364	1.28
DKFZP566A0946 protein	AL050069	1.28
integrin, alpha 2b (platelet glycoprotein IIb of IIb/IIIa complex, antigen CD41B)	AF098114	1.28
metallothionein-like 5, testis-specific (tesmin)	NM_004923	1.28
organic cation transporter OKB1	AL050350	1.28
nuclear factor of activated T-cells, cytoplasmic, calcineurin-dependent 1	U08015	1.28
gb:NM_017695.1 /DEF=Homo sapiens hypothetical protein FLJ20163 (FLJ20163), mRNA. /FEA=mRNA /GEN=FLJ20163 /PROD=hypothetical protein FLJ20163 /DB_XREF=gi:8923162 /UG=	NM_017695	1.28
proline-serine-threonine phosphatase interacting protein 2	NM_024430	1.28
hypothetical protein	AF064839	1.28
putative dipeptidase	NM_022355	1.28
apoptosis regulator BCL-G	NM_030766	1.28
Consensus includes gb:AL031600 /DEF=Human DNA sequence from clone 390E6 on chromosome 16. Contains the 3 part of the CLCN7 gene for chloride channel 7, ESTs, a GSS and four put	AL031600	1.28
ankyrin-like with transmembrane domains 1	NM_007332	1.28
arginase, type II	NM_001172	1.28
hypothetical protein FLJ20281	NM_017742	1.28
hepatocellular carcinoma-associated antigen 59	NM_016520	1.28
small inducible cytokine subfamily C, member 2	NM_003175	1.28
ephrin-A2	NM_001405	1.28
KIAA1660 protein	AB051447	1.28
hypothetical protein FLJ20457	NM_017832	1.28
hepatocyte nuclear factor 4, alpha	X87870	1.28
solute carrier family 31 (copper transporters), member 2	NM_001860	1.28
microtubule-associated protein tau	AI056359	1.28
similar to rat integral membrane glycoprotein POM121	NM_014348	1.28
indoleamine-pyrrole 2,3 dioxygenase	M34455	1.28
phosphorylase, glycogen; liver (Hers disease, glycogen storage disease type VI)	NM_002863	1.28
hypothetical protein FLJ21918	NM_024939	1.28
hypothetical protein FLJ12750	BC005882	1.28
NICE-1 protein	NM_019060	1.28
H4 histone family, member H	NM_003543	1.28
chorionic somatomammotropin hormone-like 1	NM_001318	1.28

aquaporin 5	BF726531	1.28
KIAA0255 gene product	AL515964	1.28
gb:NM_005405.1 /DEF=Homo sapiens RNA binding motif protein, Y chromosome, family 2, member B (RBMY2B) mRNA. /FEA=mRNA /GEN=RBMY2B /PROD=RNA binding motif protein, Y cl	NM_005405	1.28
meningioma (disrupted in balanced translocation) 1	NM_002430	1.28
FICSO6 binding protein 18 (12.6 kD)	NM_004116	1.28
glycine receptor, alpha 1 (startle disease/hyperekplexia, stiff man syndrome)	NM_000171	1.28
likely ortholog of mouse Arkadia	NM_017610	1.28
cytoskeleton-associated protein 4	AW029619	1.28
nuclear RNA export factor 2	NM_022053	1.28
hypothetical protein FLJ20216	NM_017715	1.28
Consensus includes gb:AI674138 /FEA=EST /DB_XREF=gi:4874618 /DB_XREF=est:wd08b06.x1 /CLONE=IMAGE:2327507 /UG=Hs.162247 ESTs, Highly similar to T-box-containing transcrij	AI674138	1.28
membrane-spanning 4-domains, subfamily A, member 6A	NM_022349	1.28
CA11	NM_019617	1.28
hemoglobin, alpha 2	AF105974	1.28
hypothetical protein FLJ23306	NM_024530	1.28
ADP-ribosylation factor-like 5	NM_012097	1.28
Consensus includes gb:AV661138 /FEA=EST /DB_XREF=gi:9882152 /DB_XREF=est:AV661138 /CLONE=GLCGPA11 /UG=Hs.282712 ESTs	AV661138	1.28
hypothetical protein FLJ23229	NM_025058	1.28
uroplakin 3	NM_006953	1.28
Huntingtin interacting protein H	AF161412	1.28
protein phosphatase 1, regulatory (inhibitor) subunit 12A	BE737620	1.28
inositol 1,4,5-triphosphate receptor, type 2	AB012610	1.28
AP1S-like 1	NM_006595	1.28
Consensus includes gb:AF164963.1 /DEF=Homo sapiens tumor antigen NA88-A pseudogene, complete sequence. /FEA=mRNA /DB_XREF=gi:5901726 /UG=Hs.306576 Homo sapiens tumo	AF164963	1.28
cystic fibrosis transmembrane conductance regulator, ATP-binding cassette (sub-family C, member 7)	W60595	1.28
solute carrier family 23 (nucleobase transporters), member 1	AF092511	1.28
hypothetical protein DKFZp564D0372	NM_031301	1.28
Consensus includes gb:AU148161 /FEA=EST /DB_XREF=gi:11009682 /DB_XREF=est:AU148161 /CLONE=MAMMA1002758 /UG=Hs.296720 Homo sapiens cDNA FLJ12396 fis, clone MAMM	AU148161	1.28
sorting nexin 13	AK001861	1.28
Rhesus blood group-associated glycoprotein	NM_000324	1.28
transcription factor 2, hepatic; LF-B3; variant hepatic nuclear factor	NM_006481	1.28
similar to rat nuclear ubiquitous casein kinase 2	NM_022731	1.28
KIAA1856 protein	BE552334	1.28
ribosomal protein L3-like	NM_005061	1.28
KIAA0480 gene product	AA635523	1.28
arachidonate 5-lipoxygenase	AI372974	1.28
Consensus includes gb:AW299245 /FEA=EST /DB_XREF=gi:6708922 /DB_XREF=est:zs48d07.x1 /CLONE=IMAGE:2772877 /UG=Hs.62601 Homo sapiens mRNA; cDNA DKFZp586K1318 (fro	AW299245	1.28
actinin, alpha 2	NM_001103	1.28
reticulon 4	AF320999	1.28
prostaglandin-endoperoxide synthase 1 (prostaglandin G/H synthase and cyclooxygenase)	NM_000962	1.28
Consensus includes gb:AW975638 /FEA=EST /DB_XREF=gi:8166855 /DB_XREF=est:EST387747 /UG=Hs.293490 ESTs	AW975638	1.28
hepatocellular carcinoma novel gene-3 protein	NM_016651	1.28
solute carrier family 11 (proton-coupled divalent metal ion transporters), member 1	AI346187	1.28
androgen-induced prostate proliferative shutoff associated protein	NM_015032	1.28
butyrophilin, subfamily 2, member A3	NM_024018	1.28
hypothetical protein FLJ11467	BC004228	1.28
ATPase, Cu++ transporting, beta polypeptide (Wilson disease)	NM_000053	1.28
high-mobility group protein 2-like 1	AL079310	1.28
glycoprotein Ib (platelet), alpha polypeptide	NM_000173	1.28
hypothetical protein MGC4663	NM_024514	1.28
hypothetical protein FLJ11125	NM_018331	1.28
CED-6 protein	AF200715	1.28
ectonucleotide pyrophosphatase/phosphodiesterase 2 (autotaxin)	L35594	1.28
ubiquitously transcribed tetratricopeptide repeat gene, X chromosome	NM_021140	1.28
UC28 protein	NM_021635	1.28
hypothetical protein FLJ22357 similar to epidermal growth factor receptor-related protein	NM_022450	1.28
KIAA0171 gene product	NM_014666	1.28
arginase, type II	U75667	1.28
CGI-96 protein	AL450314	1.28
Consensus includes gb:AI668610 /FEA=EST /DB_XREF=gi:4827918 /DB_XREF=est:yl56a05.s5 /CLONE=IMAGE:162224 /UG=Hs.32043 ESTs	AI668610	1.28
gb:BC006441.1 /DEF=Homo sapiens, Similar to RNA polymerase I transcription factor RRN3, clone MGC:13169, mRNA, complete cds. /FEA=mRNA /PROD=Similar to RNA polymerase I transc	BC006441	1.28
epididymal secretory protein ESP13.2	NM_030931	1.28
KIAA0053 gene product	D29642	1.28

carbonic anhydrase I	M33987	1.28
distal-less homeo box 5	NM_005221	1.28
Consensus includes gb:BF511678 /FEA=EST /DB_XREF=gi:11594976 /DB_XREF=est:UI-H-BI4-aom-c-02-0-UI.s1 /CLONE=IMAGE:3085418 /UG=Hs.128311 ESTs	BF511678	1.28
deleted in azoospermia 3	NM_020364	1.28
myeloid leukemia factor 1	NM_022443	1.28
gb:NM_017910.2 /DEF=Homo sapiens hypothetical protein FLJ20628 (FLJ20628), mRNA. /FEA=mRNA /GEN=FLJ20628 /PROD=hypothetical protein FLJ20628 /DB_XREF=gi:13435382 /FL	NM_017910	1.28
SFRS protein kinase 2	NM_003138	1.28
solute carrier family 13 (sodium-dependent dicarboxylate transporter), member 3	AW466889	1.28
acrosomal vesicle protein 1	NM_020114	1.28
64 protein	NM_021184	1.28
hypothetical protein	AW080196	1.28
hypothetical protein	AK023069	1.28
Consensus includes gb:BE515346 /FEA=EST /DB_XREF=gi:9722561 /DB_XREF=est:601235986F1 /CLONE=IMAGE:3608205 /UG=Hs.278871 ESTs, Weakly similar to S47072 finger protein	BE515346	1.28
tarhalin 1	NM_020127	1.28
olfactory receptor, family 11, subfamily A, member 1	NM_013937	1.28
phosphodiesterase 4B, cAMP-specific (phosphodiesterase E4 dunce homolog, Drosophila)	L20966	1.28
Consensus includes gb:AL049442.1 /DEF=Homo sapiens mRNA; cDNA DKFZp586N1720 (from clone DKFZp586N1720). /FEA=mRNA /DB_XREF=gi:4500222 /UG=Hs.71616 Homo sapiens r	AL049442	1.28
OLF-1/EBF associated zinc finger gene	AW149417	1.28
hypothetical protein FLJ21019	NM_024927	1.28
conserved gene amplified in osteosarcoma	AF022231	1.28
polycystic kidney disease 2-like 2	NM_014386	1.28
a disintegrin and metalloproteinase domain 7	AF215824	1.28
small nuclear RNA activating complex, polypeptide 2, 45kD	NM_003083	1.28
hypothetical protein FLJ20128	NM_017679	1.28
calcium-sensing receptor (hypocalciuric hypercalcemia 1, severe neonatal hyperparathyroidism)	U20760	1.28
ATP-binding cassette, sub-family B (MDR/TAP), member 1	AF016535	1.28
Consensus includes gb:AW517686 /FEA=EST /DB_XREF=gi:7155768 /DB_XREF=est:xs86d05.x1 /CLONE=IMAGE:2776521 /UG=Hs.305923 ATPase, Ca++ transporting, plasma membrane -	AW517686	1.28
runt-related transcription factor 1 (acute myeloid leukemia 1; aml1 oncogene)	AF312387	1.28
hypothetical protein FLJ23263	NM_025115	1.28
Consensus includes gb:AL133262 /DEF=Human DNA sequence from clone RP1-319D22 on chromosome 6 Contains part of the gene for a CDC5-like protein, the gene for a novel protein (PR	AL133262	1.28
zinc finger protein 261	NM_005096	1.28
trinucleotide repeat containing 11 (THR-associated protein, 230 kDa subunit)	NM_005120	1.28
DKFZP434A043 protein	BF195973	1.28
DKFZP564M1416 protein	BF981643	1.28
chromosome 21 open reading frame 108	AI803485	1.28
hypothetical protein FLJ10671	NM_018175	1.28
transient receptor potential channel 4	NM_016179	1.28
Notch homolog 2 (Drosophila)	AF308601	1.28
latent transforming growth factor beta binding protein 2	NM_000428	1.28
hypothetical protein FLJ20707	NM_017936	1.28
SEC22 vesicle trafficking protein-like 1 (S. cerevisiae)	AV701283	1.28
KIAA1616 protein	AB046836	1.28
Consensus includes gb:AA768909 /FEA=EST /DB_XREF=gi:2820147 /DB_XREF=est:nz82h01.s1 /CLONE=IMAGE:1302001 /UG=Hs.293205 ESTs, Weakly similar to BC39498 1 H.sapiens	AA768909	1.28
J domain containing protein 1	NM_021800	1.28
Dnal (Hsp40) homolog, subfamily A, member 1	AL534104	1.28
Consensus includes gb:AL121994 /DEF=Human DNA sequence from clone RP4-781L3 on chromosome 1p34.3-36.11 Contains a pseudogene similar to IFITM3 (interferon inducedntransmem	AL121994	1.28
ets variant gene 5 (ets-related molecule)	NM_004454	1.28
hypothetical protein	NM_018418	1.27
Fc fragment of IgG, low affinity IIb, receptor for (CD32)	M31933	1.27
proteoglycan 2, bone marrow (natural killer cell activator, eosinophil granule major basic protein)	BC005929	1.27
sodium channel, voltage-gated, type I, beta polypeptide	NM_001037	1.27
phosphodiesterase 4D, cAMP-specific (phosphodiesterase E3 dunce homolog, Drosophila)	U50157	1.27
hypothetical protein FLJ13310	NM_025118	1.27
Cas-Br-M (murine) ectropic retroviral transforming sequence b	NM_004351	1.27
olfactory receptor, family 5, subfamily I, member 1	NM_006637	1.27
adaptor-related protein complex 1, gamma 2 subunit	AI741833	1.27
Consensus includes gb:AL049997.1 /DEF=Homo sapiens mRNA; cDNA DKFZp564L102 (from clone DKFZp564L102). /FEA=mRNA /DB_XREF=gi:4884250 /UG=Hs.306305 Homo sapiens mR	AL049997	1.27
ATP-binding cassette, sub-family A (ABC1), member 1	NM_005502	1.27
PR domain containing 13	NM_021620	1.27
SRY (sex determining region Y)-box 11	AI360875	1.27
centaurin, delta 1	AB011152	1.27
zinc finger protein 73 (Cos12)	NM_012480	1.27
platelet-derived growth factor receptor, alpha polypeptide	H79306	1.27

hypothetical protein similar to C67943	AL138752	1.27
chromosome 9 open reading frame 9	NM_018956	1.27
hypothetical protein DKFZp566A1524	NM_030797	1.27
corneodesmosin	L20815	1.27
olfactory receptor, family 7, subfamily A, member 17	AC005255	1.27
Fc fragment of IgA, receptor for	U56237	1.27
amyloid beta (A4) precursor protein-binding, family A, member 3 (X11-like 2)	AI141541	1.27
megakaryocyte-enhanced gene transcript 1 protein	NM_021246	1.27
hypothetical protein FLJ20651	NM_017919	1.27
Consensus includes gb:AL583687 /FEA=EST /DB_XREF=gi:12952889 /DB_XREF=est:AL583687 /CLONE=CS0DI008YL09 (5 prime) /UG=Hs.134852 ESTs	AL583687	1.27
Consensus includes gb:AF052100.1 /DEF=Homo sapiens clone 23645 mRNA sequence. /FEA=mRNA /DB_XREF=gi:3360407 /UG=Hs.6651 Homo sapiens clone 23645 mRNA sequence	AF052100	1.27
ecotropic viral integration site 1	S82592	1.27
dual-specificity tyrosine-(Y)-phosphorylation regulated kinase 3	AF186773	1.27
loss of heterozygosity, 11, chromosomal region 2, gene A	BC001234	1.27
single-strand selective monofunctional uracil DNA glycosylase	NM_014311	1.27
iduronidase, alpha-L-	AI424441	1.27
myosin VIIA (Usher syndrome 1B (autosomal recessive, severe))	NM_000260	1.27
cytochrome P450, subfamily IVF, polypeptide 11	NM_021187	1.27
ornithine carbamoyltransferase	NM_000531	1.27
Consensus includes gb:AW971983 /FEA=EST /DB_XREF=gi:8161949 /DB_XREF=est:EST384192 /UG=Hs.293003 ESTs, Moderately similar to ALU4_HUMAN ALU SUBFAMILY SB2 SEQUENCE	AW971983	1.27
solute carrier family 12 (potassium/chloride transporters), member 4	U55054	1.27
guanylate cyclase 2F, retinal	NM_001522	1.27
neural cell adhesion molecule 1	AA126505	1.27
sperm protein associated with the nucleus, X chromosome, family member A1	NM_013453	1.27
leukemia inhibitory factor receptor	NM_002310	1.27
alpha-1,4-N-acetylglucosaminyltransferase	NM_016161	1.27
prostaglandin D2 synthase (21kD, brain)	M61900	1.27
ADAM-like, decysin 1	NM_014479	1.27
FSHD region gene 1	NM_004477	1.27
pleiomorphic adenoma gene-like 1	NM_006718	1.27
NCK adaptor protein 2	BC000103	1.27
secretin receptor	U13989	1.27
vasodilator-stimulated phosphoprotein	NM_003370	1.27
lectin, galactoside-binding, soluble, 8 (galectin 8)	AF342815	1.27
v-maf musculoaponeurotic fibrosarcoma oncogene homolog (avian)	NM_005360	1.27
KIAA0913 protein	W89120	1.27
activin A receptor type II-like 1	L17075	1.27
solute carrier family 17 (sodium phosphate), member 1	NM_005074	1.27
procollagen-lysine, 2-oxoglutarate 5-dioxygenase (lysine hydroxylase) 2	NM_000935	1.27
5-hydroxytryptamine (serotonin) receptor 2B	NM_000867	1.27
hypothetical protein FLJ20280	NM_017741	1.27
solute carrier family 4, sodium bicarbonate cotransporter-like, member 10	NM_022058	1.27
LIM domain kinase 2	AC002073	1.27
angiopoietin-like 3	NM_014495	1.27
hypothetical protein, expressed in osteoblast	NM_006820	1.27
carcinoembryonic antigen-related cell adhesion molecule 3	L00692	1.27
hypothetical protein FLJ13501	NM_025007	1.27
TP53TG3 protein	NM_015369	1.27
adrenergic, beta, receptor kinase 1	U08438	1.27
hypothetical protein FLJ22938	NM_024676	1.27
neuronal protein	NM_013259	1.27
epsilon-trimethyllysine hydroxylase	NM_018196	1.27
protein C receptor, endothelial (EPCR)	NM_006404	1.27
protein kinase, cAMP-dependent, regulatory, type I, alpha (tissue specific extinguisher 1)	X73110	1.27
delta-like homolog (Drosophila)	U15979	1.27
Consensus includes gb:AW954107 /FEA=EST /DB_XREF=gi:8143790 /DB_XREF=est:EST366177 /UG=Hs.303394 Homo sapiens cDNA: FLJ22150 fis, clone HRC00109	AW954107	1.27
UDP glycosyltransferase 2 family, polypeptide B15	U06641	1.27
hypothetical protein FLJ22404	NM_025043	1.27
hypothetical protein FLJ11743	NM_024527	1.27
hypothetical gene FLJ00060	AK024467	1.27
FLJ23 gene product	NM_006700	1.27
a disintegrin and metalloproteinase domain 10	N51370	1.27
hypothetical protein IMAGE 109914	NM_022076	1.27

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Consensus includes gb:AU147698 /FEA=EST /DB_XREF=gi:11009219 /DB_XREF=est:AU147698 /CLONE=MAMMA1001408 /UG=Hs.306633 Homo sapiens cDNA FLJ12248 fis, clone MAMM	AU147698	1.27
fucose-1-phosphate guanylyltransferase	NM_003838	1.27
meprin A, alpha (PABA peptide hydrolase)	NM_005588	1.27
cytochrome P450, subfamily IVB, polypeptide 1	J02871	1.27
lysosomal apyrase-like 1	AB002390	1.27
KIAA0672 gene product	NM_014859	1.27
ATPase, Ca++ transporting, plasma membrane 4	NM_001684	1.27
gamma-aminobutyric acid (GABA) A receptor, epsilon	NM_004961	1.27
steroid-5-alpha-reductase, alpha polypeptide 1 (3-oxo-5 alpha-steroid delta 4-dehydrogenase alpha 1)	NM_001047	1.27
MKP-1 like protein tyrosine phosphatase	NM_007026	1.27
suppressor of Ty 3 homolog (S. cerevisiae)	NM_003599	1.27
hypothetical protein FLJ11106	NM_018324	1.27
UDP-Gal:betaGlcNAc beta 1,3-galactosyltransferase, polypeptide 3	AB050856	1.27
Consensus includes gb:AK024679.1 /DEF=Homo sapiens cDNA: FLJ21026 fis, clone CAE06812. /FEA=mRNA /DB_XREF=gi:10437019 /UG=Hs.317585 Homo sapiens cDNA: FLJ21026 fis, cl	AK024679	1.27
clone 1900 unknown protein	NM_015883	1.27
calcium/calmodulin-dependent protein kinase (CaM kinase) II beta	AF081924	1.27
Fc fragment of IgG, low affinity IIa, receptor for (CD32)	NM_021642	1.27
microtubule-associated protein 4	AJ553791	1.27
myosin, light polypeptide 5, regulatory	L03785	1.27
neuropeptide FF-amide peptide precursor	AW195837	1.27
DKFZP564M1462 protein	AL096716	1.27
Cluster Incl. AA225490:nc25b01.s1 Homo sapiens cDNA /clone=IMAGE-1009129 /gb=AA225490 /gi=1846817 /ug=Hs.80115 /len=539	AA225490	1.27
olfactory receptor, family 7, subfamily E, member 2 pseudogene	AF065857	1.27
putative DNA/chromatin binding motif	AF087481	1.27
hypothetical protein FLJ20871 similar to FSP27	NM_022094	1.27
hypothetical protein FLJ21195 similar to protein related to DAC and cerberus	NM_022469	1.27
RAB30, member RAS oncogene family	NM_004283	1.27
gap junction protein, beta 1, 32kD (connexin 32, Charcot-Marie-Tooth neuropathy, X-linked)	NM_000166	1.27
hypothetical protein FLJ13520	NM_030896	1.27
myristoylated alanine-rich protein kinase C substrate	AW163148	1.27
gastric intrinsic factor (vitamin B synthesis)	NM_005142	1.27
corneodesmosin	NM_001264	1.27
chromosome 12 open reading frame 22	NM_030809	1.27
TONDU	BE542323	1.27
agrin	AW008051	1.27
Consensus includes gb:BG485135 /FEA=EST /DB_XREF=gi:13417414 /DB_XREF=est:602503756F1 /CLONE=IMAGE:4617445 /UG=Hs.325722 immunoglobulin kappa variable 3D-15	BG485135	1.27
interleukin 17E	NM_022789	1.27
hypothetical protein FLJ10498	NM_018115	1.27
cysteine sulfinic acid decarboxylase-related protein 2	NM_015989	1.27
PHD finger protein 1	AL021366	1.27
hypothetical protein PP1628	NM_025201	1.27
hypothetical protein FLJ11222	NM_018365	1.27
histone deacetylase 9	NM_014707	1.27
adenylate cyclase 1 (brain)	L05500	1.27
PRO1578 protein	NM_014093	1.27
Consensus includes gb:AK024561.1 /DEF=Homo sapiens cDNA: FLJ20908 fis, clone ADSE00417. /FEA=mRNA /DB_XREF=gi:10436870 /UG=Hs.306689 Homo sapiens cDNA: FLJ20908 fis, c	AK024561	1.27
Hr44 antigen	X91103	1.27
EGF-like module-containing mucin-like receptor EMR3	AF239764	1.27
UDP glycosyltransferase 1 family, polypeptide A3	NM_019093	1.27
prolactin receptor	AF349939	1.27
nitric oxide synthase 3 (endothelial cell)	NM_000603	1.27
WNT1 inducible signaling pathway protein 1	AB034725	1.27
M10098 Human 18S rRNA gene, complete (_5, _M, _3 represent transcript regions 5 prime, Middle, and 3 prime respectively)	M10098	1.27
hypothetical protein FLJ00001	AK024432	1.27
hypothetical protein DKFZp434L0850	NM_017558	1.27
tissue factor pathway inhibitor (lipoprotein-associated coagulation inhibitor)	AF021834	1.27
plasma glutamate carboxypeptidase	NM_016134	1.27
BH3 interacting domain death agonist	NM_001196	1.27
hypothetical protein FLJ20156	NM_017691	1.27
solute carrier family 6 (neurotransmitter transporter, GABA), member 1	AI003579	1.27
B1 for mucin	X83412	1.27
Consensus includes gb:AI732802 /FEA=EST /DB_XREF=gi:5053915 /DB_XREF=est:ab23e08.x5 /CLONE=IMAGE:841670 /UG=Hs.292679 ESTs	AI732802	1.27
nuclear receptor subfamily 6, group A, member 1	X99975	1.27

Consensus includes gb:Z69744 /DEF=H.sapiens ALL-1 gene exon 1 (and joined coding region) /FEA=CDS /DB_XREF=gi:1490270 /UG=Hs.199160 myeloidlymphoid or mixed-lineage leuke	Z69744	1.27
KIAA0685 gene product	AW593269	1.27
Consensus includes gb:AB028239 /DEF=Homo sapiens pseudogene for necdin /FEA=CDS /DB_XREF=gi:5672690 /UG=Hs.247843 Homo sapiens pseudogene for necdin	AB028239	1.27
aryl hydrocarbon receptor nuclear translocator	AL042842	1.27
spondin 1, (f-spondin) extracellular matrix protein	AB018305	1.27
immunoglobulin heavy constant mu	U80164	1.27
calcium channel, voltage-dependent, gamma subunit 1	NM_000727	1.27
thousand and one amino acid protein kinase	NM_004783	1.27
paired box gene 4	AB008913	1.27
Consensus includes gb:AI129381 /FEA=EST /DB_XREF=gi:3597895 /DB_XREF=est:qc34d02.x1 /CLONE=IMAGE:1711491 /UG=Hs.89463 potassium large conductance calcium-activated ch	AI129381	1.27
Consensus includes gb:AI471375 /FEA=EST /DB_XREF=gi:4333465 /DB_XREF=est:tm10c12.x1 /CLONE=IMAGE:2156182 /UG=Hs.80206 glucose-6-phosphate dehydrogenase	AI471375	1.27
activin A receptor, type IB	NM_020327	1.27
chromobox homolog 2 (Pc class homolog, Drosophila)	BE258133	1.27
Consensus includes gb:AU146949 /FEA=EST /DB_XREF=gi:11008470 /DB_XREF=est:AU146949 /CLONE=HEMBB1001952 /UG=Hs.301817 Homo sapiens cDNA FLJ12044 fis, clone HEMBB	AU146949	1.27
histone acetyltransferase	NM_012330	1.27
hypothetical protein FLJ20429	NM_017817	1.27
KIAA1446 protein	NM_020836	1.27
Consensus includes gb:AI967961 /FEA=EST /DB_XREF=gi:5764779 /DB_XREF=est:wu11d08.x1 /CLONE=IMAGE:2516655 /UG=Hs.26837 Homo sapiens mRNA; cDNA DKFZp586K1123 (fro	AI967961	1.27
protein kinase C-like 2	BG292065	1.27
T-cell lymphoma invasion and metastasis 2	NM_012454	1.27
gb:BC001279.1 /DEF=Homo sapiens, Similar to KIAA0626 gene product, clone MGC:5129, mRNA, complete cds. /FEA=mRNA /PROD=Similar to KIAA0626 gene product /DB_XREF=gi:12654	BC001279	1.27
UDP-GlcNAc:betaGal beta-1,3-N-acetylglucosaminyltransferase 4	NM_030765	1.27
heterogeneous nuclear ribonucleoprotein H1 (H)	NM_005520	1.27
gb:U03891.2 /DEF=Homo sapiens phorbolin I mRNA, complete cds. /FEA=mRNA /PROD=phorbolin I /DB_XREF=gi:4895107 /UG=Hs.226307 phorbolin (similar to apolipoprotein B mRNA e	U03891	1.27
hypothetical protein MGC3146	NM_031289	1.27
tissue inhibitor of metalloproteinase 3 (Sorsby fundus dystrophy, pseudoinflammatory)	U67195	1.27
Consensus includes gb:AL122093.1 /DEF=Homo sapiens mRNA; cDNA DKFZp43482115 (from clone DKFZp43482115). /FEA=mRNA /DB_XREF=gi:6102904 /UG=Hs.136241 Homo sapiens	AL122093	1.27
hypothetical protein FLJ13639	AW194384	1.27
protease, serine, 8 (prostasin)	NM_002773	1.27
Consensus includes gb:AA224446 /FEA=EST /DB_XREF=gi:1844989 /DB_XREF=est:zr16a03.r1 /CLONE=IMAGE:663532 /UG=Hs.311561 ESTs, Weakly similar to CYA7_HUMAN ADENYLATE	AA224446	1.27
integrin, beta 1 (fibronectin receptor, beta polypeptide, antigen CD29 includes MDF2, MSK12)	BG435463	1.27
caspase 2, apoptosis-related cysteine protease (neural precursor cell expressed, developmentally down-regulated 2)	U13022	1.27
osteoblast specific factor 2 (fasciclin I-like)	D13665	1.27
pregnancy specific beta-1-glycoprotein 6	NM_002782	1.27
Cluster Incl. AI557319:PT2.1_16_F08.r Homo sapiens cDNA, 3 end /clone_end=3 /gb=AI557319 /gi=4489682 /ug=Hs.21921 /len=876	AI557319	1.27
SRY (sex determining region Y)-box 9 (campomelic dysplasia, autosomal sex-reversal)	AI382146	1.27
desmuslin	AK026420	1.27
hypothetical protein PRO1097	NM_018499	1.27
Consensus includes gb:AK025114.1 /DEF=Homo sapiens cDNA: FLJ21461 fis, clone COL04727. /FEA=mRNA /DB_XREF=gi:10437566 /UG=Hs.306768 Homo sapiens cDNA: FLJ21461 fis, cl	AK025114	1.27
hypothetical protein FLJ22501	NM_024747	1.27
Consensus includes gb:AL038824 /FEA=EST /DB_XREF=gi:5407955 /DB_XREF=est:DKFZp56602446_s1 /CLONE=DKFZp56602446 /UG=Hs.12840 Homo sapiens germline mRNA sequenc	AL038824	1.27
hypothetical protein	AL157485	1.27
ciliary neurotrophic factor	NM_000614	1.27
tripartite motif-containing 14	BE962483	1.27
aquaporin 8	NM_001169	1.27
docking protein 1, 62kD (downstream of tyrosine kinase 1)	AF035299	1.27
protein tyrosine phosphatase, non-receptor type 21	NM_007039	1.27
copine VII	NM_014427	1.27
mucin 2, intestinal/tracheal	NM_002457	1.27
Consensus includes gb:AA664011 /FEA=EST /DB_XREF=gi:2618002 /DB_XREF=est:ac03f10.s1 /CLONE=IMAGE:855403 /UG=Hs.173638 transcription factor 7-like 2 (T-cell specific, HMG-b	AA664011	1.27
topoisomerase (DNA) II binding protein	NM_007027	1.27
COP9 homolog	BC003090	1.27
complement component (3b/4b) receptor 1, including Knops blood group system	NM_000573	1.27
ARS component B	NM_020427	1.27
aquaporin 4	U63622	1.27
Nisshapen/NIK-related kinase	AF218033	1.27
brefeldin A-inhibited guanine nucleotide-exchange protein 2	AV657604	1.27
Fc fragment of IgG, low affinity IIIb, receptor for (CD16)	NM_000570	1.27
IK cytokine, down-regulator of HLA II	AF182645	1.27
hypoxia-inducible factor 1, alpha subunit inhibitor	NM_017902	1.27
ZW10 interactor	NM_007057	1.27
KIAA0077 protein	AU143855	1.27
HZFw1 protein	NM_025236	1.27

Consensus includes gb:AK024867.1 /DEF=Homo sapiens cDNA: FLJ21214 fis, clone COL00523. /FEA=mRNA /DB_XREF=gi:10437277 /UG=Hs.306710 Homo sapiens cDNA: FLJ21214 fis, cl	AK024867	1.27
cytochrome P450, subfamily IIC (mephenytoin 4-hydroxylase), polypeptide 9	BE971373	1.27
glutathione S-transferase M4	NM_000850	1.27
spinocerebellar ataxia 2 (olivopontocerebellar ataxia 2, autosomal dominant, ataxin 2)	NM_002973	1.27
pre-alpha (globulin) inhibitor, H3 polypeptide	NM_002217	1.27
KIAA0061 protein	D31765	1.27
CGI-78 protein	NM_016022	1.27
hypothetical protein MGC4659	NM_025268	1.27
KIAA1405 protein	AA909345	1.27
HMG-box containing protein 1	NM_012257	1.27
hypothetical protein FLJ10520	BC002574	1.27
secretory leukocyte protease inhibitor (antileukoproteinase)	NM_003064	1.27
hypothetical protein FLJ11560	AL583909	1.27
diaphanous homolog 3 (Drosophila)	NM_030932	1.27
kidney- and liver-specific gene	AB013094	1.26
Consensus includes gb:AK025422.1 /DEF=Homo sapiens cDNA: FLJ21769 fis, clone COLF7354. /FEA=mRNA /DB_XREF=gi:10437930 /UG=Hs.306810 Homo sapiens cDNA: FLJ21769 fis, cl	AK025422	1.26
sarcoglycan, delta (35kD dystrophin-associated glycoprotein)	U58331	1.26
adaptor-related protein complex 1, gamma 1 subunit	BF752277	1.26
hypothetical protein FLJ23235	NM_024943	1.26
hypothetical protein FLJ13273	NM_024751	1.26
glycoprotein A33 (transmembrane)	NM_005814	1.26
vaccinia related kinase 1	NM_003384	1.26
Consensus includes gb:AK022170.1 /DEF=Homo sapiens cDNA FLJ12108 fis, clone MAMMA1000009. /FEA=mRNA /DB_XREF=gi:10433505 /UG=Hs.250813 Homo sapiens cDNA FLJ12108 f	AK022170	1.26
f-box and WD-40 domain protein 3	AA845710	1.26
neurogenic differentiation 4	NM_021191	1.26
cryptochrome 1 (photolyase-like)	D83702	1.26
MCG/V-6 protein	AF036973	1.26
KIAA0748 gene product	NM_014796	1.26
succinate dehydrogenase complex, subunit B, iron sulfur (Ip)	AW294107	1.26
hypothetical protein PRO2176	BE552428	1.26
LIM domain binding 1	NM_003893	1.26
prostate epithelium-specific Ets transcription factor	AI307915	1.26
ribosomal protein L17	AA522618	1.26
CASP8 and FADD-like apoptosis regulator	U97075	1.26
Consensus includes gb:AK022038.1 /DEF=Homo sapiens cDNA FLJ11976 fis, clone HEM8B1001253. /FEA=mRNA /DB_XREF=gi:10433356 /UG=Hs.296679 Homo sapiens cDNA FLJ11976 f	AK022038	1.26
dachshund homolog (Drosophila)	AW772082	1.26
GATA binding protein 4	AV700724	1.26
gonadotropin-releasing hormone receptor	Z81148	1.26
tumor necrosis factor receptor superfamily, member 1B	NM_001066	1.26
FK506 binding protein 8 (38kD)	NM_012181	1.26
Consensus includes gb:AL136318 /DEF=Human DNA sequence from clone RP1-254P11 on chromosome 10 Contains ESTs, STSs and GSSs. Contains a gene for a novel protein similar to RNA	AL136318	1.26
PRO0149 protein	NM_014117	1.26
ribonuclease, RNase A family, 4	AI761728	1.26
topoisomerase (DNA) III alpha	NM_004618	1.26
Ras-GTPase-activating protein SH3-domain-binding protein	BG500067	1.26
regulator of G-protein signalling 1	NM_002922	1.26
Consensus includes gb:AA530995 /FEA=EST /DB_XREF=gi:2273701 /DB_XREF=est:nj07a02.s1 /CLONE=IMAGE:985610 /UG=Hs.23729 Homo sapiens clone 24405 mRNA sequence	AA530995	1.26
major histocompatibility complex, class II, DQ beta 1	AW276186	1.26
transcription factor 3 (E2A immunoglobulin enhancer binding factors E12/E47)	BE962186	1.26
Consensus includes gb:AL133262 /DEF=Human DNA sequence from clone RP1-319D22 on chromosome 6 Contains part of the gene for a CDC5-like protein, the gene for a novel protein (PR	AL133262	1.26
hypothetical protein FLJ20896	NM_024605	1.26
a disintegrin-like and metalloprotease (reprolysin type) with thrombospondin type 1 motif, 8	NM_007037	1.26
cullin 4B	AF212995	1.26
cytochrome P450, subfamily IIC (mephenytoin 4-hydroxylase), polypeptide 9	M21940	1.26
myosin, heavy polypeptide 7, cardiac muscle, beta	AI292276	1.26
hemochromatosis	AF144242	1.26
hypothetical protein FLJ14351	NM_024732	1.26
protein tyrosine phosphatase, receptor type, R	U77917	1.26
KIAA1061 protein	AI635302	1.26
protein kinase C, gamma	NM_002739	1.26
distal-less homeobox 4	AF254115	1.26
interferon-induced protein 35	BC001356	1.26
5-hydroxytryptamine (serotonin) receptor 2C	NM_000868	1.26

chromosome 11 open reading frame 16	NM_020643	1.26
Consensus includes gb:U79297.1 /DEF=Human clone 23589 mRNA sequence. /FEA=mRNA /DB_XREF=gi:1710280 /UG=Hs.11506 Human clone 23589 mRNA sequence	U79297	1.26
T cell receptor alpha locus	AE000659	1.26
alcohol dehydrogenase 1A (class I), alpha polypeptide	NM_000667	1.26
RAB3A interacting protein (rabin3)-like 1	NM_013401	1.26
formyl peptide receptor 1	M60626	1.26
chorionic somatomammotropin hormone-like 1	NM_022581	1.26
forkhead box O3A	N25732	1.26
Consensus includes gb:F09448 /FEA=EST /DB_XREF=gi:681981 /DB_XREF=est:HSC31G032 /CLONE=c-31g03 /UG=Hs.12840 Homo sapiens germline mRNA sequence	F09448	1.26
RAS guanyl releasing protein 2 (calcium and DAG-regulated)	AI688812	1.26
chromosome 6 open reading frame 5	AW296296	1.26
C/EBP-induced protein	NM_030802	1.26
DNA-dependent protein kinase catalytic subunit-interacting protein 2	BG251252	1.26
Consensus includes gb:AL109674.1 /DEF=Homo sapiens mRNA full length insert cDNA clone EUROIMAGE 46866. /FEA=mRNA /DB_XREF=gi:5689832 /UG=Hs.221811 Homo sapiens mRNA	AL109674	1.26
KIT ligand	AF119835	1.26
hyperpolarization activated cyclic nucleotide-gated potassium channel 2	AI421964	1.26
protein phosphatase 1B (formerly 2C), magnesium-dependent, beta isoform	AF136972	1.26
toll-like receptor 5	AF051151	1.26
Consensus includes gb:AF070610.1 /DEF=Homo sapiens clone 24505 mRNA sequence. /FEA=mRNA /DB_XREF=gi:3387992 /UG=Hs.100543 Homo sapiens clone 24505 mRNA sequence	AF070610	1.26
Misshapen/NIK-related kinase	AL157418	1.26
matrix metalloproteinase 3 (stromelysin 1, progelatinase)	NM_002422	1.26
serine dehydratase	NM_006843	1.26
small inducible cytokine subfamily A (Cys-Cys), member 18, pulmonary and activation-regulated	Y13710	1.26
Consensus includes gb:AF098968.1 /DEF=Homo sapiens familial Mediterranean fever locus region, mRNA sequence. /FEA=mRNA /DB_XREF=gi:4323318 /UG=Hs.113111 Homo sapiens fa	AF098968	1.26
brefeldin A-inhibited guanine nucleotide-exchange protein 1	NM_006421	1.26
Consensus includes gb:AU154785 /FEA=EST /DB_XREF=gi:11016306 /DB_XREF=est:AU154785 /CLONE=NT2RP4002888 /UG=Hs.31532 Homo sapiens mRNA; cDNA DKFZp434F172 (from	AU154785	1.26
homeo box D3	NM_006898	1.26
MT-protocadherin	AI825832	1.26
Consensus includes gb:AW974812 /FEA=EST /DB_XREF=gi:8166015 /DB_XREF=est:EST386917 /UG=Hs.291971 ESTs	AW974812	1.26
KIAA0685 gene product	NM_014678	1.26
hypothetical protein FLJ21687	NM_024859	1.26
actin binding LIM protein 1	BC002448	1.26
cell cycle related kinase	NM_012119	1.26
similar to S. cerevisiae RER1	AI818736	1.26
hypothetical protein FLJ22476	NM_025044	1.26
cyclin D2	NM_001759	1.26
prostate androgen-regulated transcript 1	AI770098	1.26
COQ7 coenzyme Q, 7 homolog ubiquinone (yeast)	AK024291	1.26
hypothetical protein FLJ11100	NM_018321	1.26
protein kinase C substrate 80K-H	AI815793	1.26
Consensus includes gb:BC001879.1 /DEF=Homo sapiens, clone IMAGE:3534181, mRNA, partial cds. /FEA=mRNA /PROD=Unknown (protein for IMAGE:3534181) /DB_XREF=gi:12804862 /L	BC001879	1.26
mitogen-activated protein kinase 8 interacting protein 3	AB028989	1.26
chondroitin sulfate proteoglycan BEHAB/brevican	AF229053	1.26
olfactory receptor, family 7, subfamily E, member 12 pseudogene	AF065854	1.26
myeloperoxidase	NM_000250	1.26
collagen, type IV, alpha 4	D17391	1.26
Consensus includes gb:AK021836.1 /DEF=Homo sapiens cDNA FLJ11774 fis, clone HEMBA1005884. /FEA=mRNA /DB_XREF=gi:10433104 /UG=Hs.293917 Homo sapiens cDNA FLJ11774 fi	AK021836	1.26
putative transmembrane protein	NM_012342	1.26
splicing factor 3a, subunit 1, 120kD	NM_005877	1.26
chitinase 3-like 2	U58515	1.26
BUB1 budding uninhibited by benzimidazoles 1 homolog (yeast)	AK023540	1.26
DKFZP586B2420 protein	BF509657	1.26
Consensus includes gb:AW238724 /FEA=EST /DB_XREF=gi:6571113 /DB_XREF=est:xp30h08.x1 /CLONE=IMAGE:2741919 /UG=Hs.314230 ESTs, Highly similar to clock H.sapiens	AW238724	1.26
EGF-TM7-latrophilin-related protein	NM_022159	1.26
imprinted in Prader-Willi syndrome	AI672541	1.26
polyglutamine binding protein 1	AB041834	1.26
Consensus includes gb:AK022254.1 /DEF=Homo sapiens cDNA FLJ12192 fis, clone MAMMA1000851. /FEA=mRNA /DB_XREF=gi:10433612 /UG=Hs.288451 Homo sapiens cDNA FLJ12192 f	AK022254	1.26
ring finger protein 1	AL031228	1.26
Misshapen/NIK-related kinase	AL157418	1.26
low density lipoprotein receptor-related protein 4	AAS84297	1.26
cytosolic beta-glucosidase	NM_020973	1.26
KIAA0653 protein, B7-like protein	AI659611	1.26
potassium voltage-gated channel, shaker-related subfamily, beta member 1	U33428	1.26

Consensus includes gb:AF059180 /DEF=Homo sapiens mutant beta-globin (HBB) gene, complete cds /FEA=mRNA /DB_XREF=gi:4837722 /UG=Hs.155376 hemoglobin, beta	AF059180	1.26
transglutaminase 3 (E polypeptide, protein-glutamine-gamma-glutamyltransferase)	NM_003245	1.26
hypothetical protein FLJ10665	NM_018173	1.26
angiopoietin 2	AF187858	1.26
Consensus includes gb:AJ408433 /DEF=Homo sapiens partial IGKV gene for immunoglobulin kappa chain variable region, clone 38 /FEA=CDS /DB_XREF=gi:12655549 /UG=Hs.307136 Hom	AJ408433	1.26
retinol binding protein 4, plasma	AF119868	1.26
chromosome 2 open reading frame 6	NM_018221	1.26
sushi-repeat protein	NM_014467	1.26
chromosome 11open reading frame 9	NM_013279	1.26
Consensus includes gb:BE466926 /FEA=EST /DB_XREF=gi:9512701 /DB_XREF=est:hz59a04.x1 /CLONE=IMAGE:3212238 /UG=Hs.279706 ESTs	BE466926	1.26
interleukin 13 receptor, alpha 1	U81380	1.26
Consensus includes gb:BG236220 /FEA=EST /DB_XREF=gi:12750151 /DB_XREF=est:naf25h11.x1 /CLONE=IMAGE:4142085 /UG=Hs.323067 Homo sapiens mRNA; cDNA DKFZp434I194 (h	BG236220	1.26
ATPase, Ca++ transporting, type 2C, member 1	AF225981	1.26
gb:AF165520.1 /DEF=Homo sapiens phorbolin I protein (PBI) mRNA, complete cds. /FEA=mRNA /GEN=PBI /PROD=phorbolin I protein /DB_XREF=gi:9294746 /UG=Hs.8583 similar to APOI	AF165520	1.26
binder of Arl Two	BC003087	1.26
prostaglandin E receptor 3 (subtype EP3)	D38298	1.26
synovial sarcoma, X breakpoint 2	X79200	1.26
microtubule-associated protein, RP/EB family, member 3	AI885178	1.26
gb:AF212224.1 /DEF=Homo sapiens CLK4 mRNA, complete cds. /FEA=mRNA /PROD=CLK4 /DB_XREF=gi:9437514 /UG=Hs.295231 Homo sapiens CLK4 mRNA, complete cds /FL=gb:AF212	AF212224	1.26
mitogen-activated protein kinase kinase 3	NM_002756	1.26
DKFZP434O047 protein	NM_015594	1.26
Consensus includes gb:AI889513 /FEA=EST /DB_XREF=gi:5594677 /DB_XREF=est:wn05e06.x1 /CLONE=IMAGE:2444578 /UG=Hs.168694 Homo sapiens clone 23763 unknown mRNA, part	AI889513	1.26
Consensus includes gb:AL049430.1 /DEF=Homo sapiens mRNA; cDNA DKFZp586H201 (from clone DKFZp586H201). /FEA=mRNA /DB_XREF=gi:4500210 /UG=Hs.302061 Homo sapiens mI	AL049430	1.26
trinucleotide repeat containing 21	U80756	1.26
solute carrier family 24 (sodium/potassium/calcium exchanger), member 1	NM_004727	1.26
Consensus includes gb:AK026778.1 /DEF=Homo sapiens cDNA: FLJ23125 fis, clone LNG08217. /FEA=mRNA /DB_XREF=gi:10439709 /UG=Hs.76325 step II splicing factor SLU7	AK026778	1.26
Rhesus blood group, D antigen	X63097	1.26
ATPase, aminophospholipid transporter-like, Class I, type 8A, member 2	AU146927	1.26
N-acetylglucosaminidase, alpha- (Sanfilippo disease IIIB)	AI364950	1.26
serine/threonine kinase 17a (apoptosis-inducing)	AW183478	1.26
zinc finger protein 274	NM_016325	1.26
EphA4	NM_004438	1.26
myelin-associated oligodendrocyte basic protein	NM_006501	1.26
tousled-like kinase 1	NM_012290	1.26
hypothetical protein DKFZp762P2111	R61320	1.26
peroxisome proliferative activated receptor, delta	L07592	1.26
Consensus includes gb:AU146050 /FEA=EST /DB_XREF=gi:11007571 /DB_XREF=est:AU146050 /CLONE=HEMBA1006665 /UG=Hs.287459 Homo sapiens cDNA FLJ11844 fis, clone HEMBA	AU146050	1.26
hypothetical protein FLJ13386	NM_025180	1.26
platelet-derived growth factor beta polypeptide (simian sarcoma viral (v-sis) oncogene homolog)	NM_002608	1.26
matrix metalloproteinase 20 (enamelysin)	NM_004771	1.26
Consensus includes gb:AL046350 /FEA=EST /DB_XREF=gi:5434427 /DB_XREF=est:DKFZp434J097_s1 /CLONE=DKFZp434J097 /UG=Hs.225433 Homo sapiens mRNA; cDNA DKFZp434G15	AL046350	1.26
5,10-methenyltetrahydrofolate synthetase (5-formyltetrahydrofolate cyclo-ligase)	NM_006441	1.26
carboxypeptidase Z	BC006393	1.26
a disintegrin-like and metalloprotease (reprolysin type) with thrombospondin type 1 motif, 7	NM_014272	1.26
Consensus includes gb:AK025064.1 /DEF=Homo sapiens cDNA: FLJ21411 fis, clone COL03986. /FEA=mRNA /DB_XREF=gi:10437503 /UG=Hs.306758 Homo sapiens cDNA: FLJ21411 fis, cl	AK025064	1.26
stromal interaction molecule 1	NM_003156	1.26
ATP-binding cassette, sub-family B (MDR/TAP), member 6	NM_005689	1.26
retinoid X receptor, alpha	BE675800	1.26
similar to proline-rich protein 48	NM_019043	1.26
Cluster Incl. T87245:yd91b02.r1 Homo sapiens cDNA, 5' end /clone=IMAGE-115563 /clone_end=5' /gb=T87245 /gi=715597 /ug=Hs.16336 /len=412	T87245	1.26
melanoma antigen, family A, 12	BC003408	1.26
hypothetical protein MGC5306	NM_024116	1.26
hyaluronan-mediated motility receptor (RHAMM)	NM_012485	1.26
Consensus includes gb:BF347113 /FEA=EST /DB_XREF=gi:11294708 /DB_XREF=est:602020879F1 /CLONE=IMAGE:4156602 /UG=Hs.273138 ESTs	BF347113	1.26
protein kinase H11	AF133207	1.26
potassium inwardly-rectifying channel, subfamily J, member 10	NM_002241	1.26
dedicator of cyto-kinesis 3	BF593175	1.26
pyruvate dehydrogenase kinase, isoenzyme 2	AL574319	1.26
organic cationic transporter-like 4	NM_004803	1.26
glycophorin B (includes Ss blood group)	NM_002100	1.26
EGF-like-domain, multiple 6	NM_015507	1.26
v-ets musculoaponeurotic fibrosarcoma oncogene homolog F (avian)	AA725102	1.26
Consensus includes gb:AW340588 /FEA=EST /DB_XREF=gi:6837214 /DB_XREF=est:hd05a05.x1 /CLONE=IMAGE:2908592 /UG=Hs.8736 Homo sapiens mRNA; cDNA DKFZp564H203 (from	AW340588	1.26

aldolase B, fructose-bisphosphate	AK026411	1.26
ATP-binding cassette, sub-family B (MDR/TAP), member 9	NM_019625	1.26
dystrobrevin, beta	AK022277	1.26
ceroid-lipofuscinosis, neuronal 8 (epilepsy, progressive with mental retardation)	AF123759	1.26
hypothetical protein FLJ14297	NM_024903	1.26
hypothetical protein FLJ10698	AL050219	1.26
guanine nucleotide binding protein (G protein) alpha 12	AK024696	1.26
brain cell membrane protein 1	AL136550	1.26
KIAA1069 protein	AW665865	1.26
WNT1 inducible signaling pathway protein 1	NM_003882	1.26
ficolin (collagen/fibrinogen domain-containing) 3 (Hakata antigen)	NM_003665	1.26
hypothetical protein FLJ10738	NM_018199	1.26
interferon-induced protein with tetratricopeptide repeats 4	NM_001549	1.26
carbonyl reductase 3	NM_001236	1.26
class I cytokine receptor	NM_004843	1.26
carbohydrate (chondroitin 6/keratan) sulfotransferase 3	NM_004273	1.26
IMB (bacterial acetolactate synthase)-like	BC000109	1.26
glycine receptor, beta	AF094754	1.26
homeo box B13	U57052	1.26
KIAA1624 protein	NM_020960	1.26
Consensus includes gb:AF229166.1 /DEF=Homo sapiens docking protein 1-like protein mRNA, partial cds. /FEA=mRNA /PROD=docking protein 1-like protein /DB_XREF=gi:7650361 /UG=H	AF229166	1.26
S100 calcium binding protein A6 (calcyclin)	NM_014624	1.26
thyroid hormone receptor interactor 13	NM_004237	1.26
small nuclear ribonucleoprotein polypeptide F	NM_003095	1.26
homeo box B7	S49765	1.26
hypothetical protein FLJ10713	NM_018189	1.26
stromal cell-derived factor 1	NM_000609	1.26
unc93 (C.elegans) homolog B	NM_030930	1.26
growth associated protein 43	AF279774	1.26
KIAA0844 protein	NM_014951	1.26
Consensus includes gb:AV759552 /FEA=EST /DB_XREF=gi:10917400 /DB_XREF=est:AV759552 /CLONE=MDSCBH06 /UG=Hs.226138 Homo sapiens mRNA; cDNA DKFZp566H2446 (from cl	AV759552	1.26
PDZ domain containing 1	NM_002614	1.26
Putative prostate cancer tumor suppressor	U42349	1.26
receptor (calcitonin) activity modifying protein 3	NM_005856	1.26
NCK adaptor protein 1	BC006403	1.26
hypothetical protein MGC2849	NM_024080	1.26
hypothetical protein DKFZp566H0824	NM_017535	1.26
Consensus includes gb:AF090887.1 /DEF=Homo sapiens clone HQ0085. /FEA=mRNA /DB_XREF=gi:6690154 /UG=Hs.306562 Homo sapiens clone HQ0085	AF090887	1.26
hypothetical protein FLJ11082	NM_018317	1.26
Consensus includes gb:AF103295.1 /DEF=Homo sapiens clone N97 immunoglobulin heavy chain variable region mRNA, partial cds. /FEA=mRNA /PROD=immunoglobulin heavy chain variabl	AF103295	1.26
Cluster Incl. AL109698:Homo sapiens mRNA full length insert cDNA clone EUROIMAGE 26539 /cds=UNKNOWN /gb=AL109698 /gi=5689808 /ug=Hs.8065 /len=2035	AL109698	1.26
guanine nucleotide binding protein (G protein), alpha 14	NM_004297	1.26
vacuolar protein sorting 33B (yeast)	NM_018668	1.26
Consensus includes gb:AL133269 /DEF=Human DNA sequence from clone RP3-468K3 on chromosome 6q25-26. Contains a beta tubulin pseudogene, STSs and GSSs /FEA=CDS /DB_XREF=;	AL133269	1.26
stomatin (EPB72)-like 1	Y16522	1.26
hypothetical protein FLJ21472	NM_024837	1.26
forkhead box G1B	NM_005249	1.26
Consensus includes gb:AL050148.1 /DEF=Homo sapiens mRNA; cDNA DKFZp586G1520 (from clone DKFZp586G1520). /FEA=mRNA /DB_XREF=gi:4884359 /UG=Hs.31834 Homo sapiens c	AL050148	1.26
tumor necrosis factor (TNF superfamily, member 2)	NM_000594	1.26
gb:NM_016528.1 /DEF=Homo sapiens hydroxyacid oxidase 3 (medium-chain) (HAO3), mRNA. /FEA=mRNA /GEN=HAO3 /PROD=hydroxyacid oxidase 3 /DB_XREF=gi:7706074 /FL=gb:NM_	NM_016528	1.26
likely homolog of rat and mouse retinoid-inducible serine carboxypeptidase	NM_021626	1.26
cartilage oligomeric matrix protein (pseudoachondroplasia, epiphyseal dysplasia 1, multiple)	NM_000095	1.26
protocadherin 12	NM_016580	1.26
protocadherin gamma subfamily A, 9	AF152516	1.26
hypothetical protein DKFZp434I143	AL080135	1.26
Fc fragment of IgA, receptor for	D87858	1.25
ephrin-B3	NM_001406	1.25
casein, beta	NM_001891	1.25
Rho-associated, coiled-coil containing protein kinase 1	N22548	1.25
fem-1 homolog b (C. elegans)	NM_015322	1.25
interleukin 5 receptor, alpha	M75914	1.25
protein kinase NYD-SP15	NM_030911	1.25
leukocyte immunoglobulin-like receptor, subfamily B (with TM and ITIM domains), member 2	AF004231	1.25

QP 5

type 1 tumor necrosis factor receptor shedding aminopeptidase regulator	A8011097	1.47
hypothetical protein FLJ12929	NM_024937	1.47
azoospermia factor 1	AC007320	1.47
hypothetical protein FLJ13397	NM_024948	1.47
mitochondrial ribosomal protein 63	NM_024026	1.47
solute carrier family 13 (sodium/sulfate symporters), member 1	AF260824	1.47
zinc finger protein 297	AL523144	1.47
Consensus includes gb:AF052173.1 /DEF=Homo sapiens clone 24417 mRNA sequence. /FEA=mRNA /DB_XREF=gi:3360484 /UG=Hs.306253 Homo sapiens clone 24417 mRNA sequence	AF052173	1.47
KIAA1035 protein	AB028958	1.47
protein Z, vitamin K-dependent plasma glycoprotein	NM_003891	1.47
purinergic receptor P2X-like 1, orphan receptor	AF065385	1.47
bone morphogenetic protein receptor, type II (serine/threonine kinase)	U25110	1.47
tumor necrosis factor (ligand) superfamily, member 11	AF053712	1.47
estrogen-related receptor gamma	NM_001438	1.47
leukotriene C4 synthase	NM_000897	1.47
mitochondrial ribosomal protein L9	AW611641	1.47
NADPH-dependent FMN and FAD containing oxidoreductase	NM_014434	1.47
ras homolog gene family, member I	AK021882	1.47
SIX-like	X15217	1.47
KIAA1649 protein	AB055760	1.47
Consensus includes gb:BF508835 /FEA=EST /DB_XREF=gi:11592133 /DB_XREF=est:UI-H-BI4-aor-d-03-0-UI.s1 /CLONE=IMAGE:3085853 /UG=Hs.6217 Homo sapiens cDNA FLJ12521 fis,	BF508835	1.47
Consensus includes gb:AI806793 /FEA=EST /DB_XREF=gi:5393359 /DB_XREF=est:wf15d05.x1 /CLONE=IMAGE:2350665 /UG=Hs.249239 collagen, type VIII, alpha 2	AI806793	1.47
tumor-associated calcium signal transducer 2	NM_002353	1.47
crystallin, mu	NM_001888	1.47
Purkinje cell protein 4	NM_006198	1.47
HEAT-like repeat-containing protein	NM_020958	1.47
nuclear receptor subfamily 4, group A, member 3	NM_006981	1.47
frizzled homolog 10 (Drosophila)	NM_007197	1.47
capping protein (actin filament) muscle Z-line, beta	BG420053	1.47
fetuin B	N39010	1.47
WW domain-containing oxidoreductase	U13395	1.47
Consensus includes gb:AI086530 /FEA=EST /DB_XREF=gi:3424953 /DB_XREF=est:oz72a04.x1 /CLONE=IMAGE:1680846 /UG=Hs.255883 ESTs	AI086530	1.47
a disintegrin and metalloproteinase domain 3a (cyritestin 1)	X89657	1.47
Consensus includes gb:AK026723.1 /DEF=Homo sapiens cDNA: FLJ23070 fis, clone LNG05629. /FEA=mRNA /DB_XREF=gi:10439641 /UG=Hs.30579 Homo sapiens cDNA: FLJ23070 fis, clo	AK026723	1.47
ring finger protein 15	NM_006355	1.47
gb:NM_018495.3 /DEF=Homo sapiens NAG22 protein (LOC55873), mRNA. /FEA=mRNA /GEN=LOC55873 /PROD=NAG22 protein /DB_XREF=gi:13236500 /UG=Hs.283080 NAG22 protein /	NM_018495	1.47
sex determining region Y	NM_003140	1.47
nucleosome assembly protein 1-like 1	BC002387	1.47
cold shock domain protein A	AW170359	1.47
hypothetical protein similar to tenascin-R	BF432086	1.47
sialic acid binding Ig-like lectin 8	AF287892	1.47
Consensus includes gb:AL079294.1 /DEF=Homo sapiens mRNA full length insert cDNA clone EUROIMAGE 362780. /FEA=mRNA /DB_XREF=gi:5102751 /UG=Hs.29952 Homo sapiens mRNA	AL079294	1.47
Consensus includes gb:AL512688.1 /DEF=Homo sapiens mRNA; cDNA DKFZp547I2313 (from clone DKFZp547I2313). /FEA=mRNA /GEN=DKFZp547I2313 /PROD=hypothetical protein /DB_	AL512688	1.47
sarcoglycan, beta (43kD dystrophin-associated glycoprotein)	NM_000232	1.47
gb:NM_007116.1 /DEF=Homo sapiens tenascin XA (TNXA), mRNA. /FEA=mRNA /GEN=TNXA /PROD=tenascin XA /DB_XREF=gi:6005907 /UG=Hs.283750 tenascin XA /FL=gb:U24488.1 gb	NM_007116	1.47
Consensus includes gb:D60132 /FEA=EST /DB_XREF=gi:961771 /DB_XREF=est:HUM087G05A /CLONE=GEN-087G05 /UG=Hs.79981 Human clone 23560 mRNA sequence	D60132	1.47
CLL7 protein	NM_018191	1.47
carbonic anhydrase VII	NM_005182	1.47
ankyrin repeat domain 2 (stretch responsive muscle)	NM_020349	1.47
Consensus includes gb:AF037219.1 /DEF=Homo sapiens PIX1 mRNA sequence. /FEA=mRNA /DB_XREF=gi:4090971 /UG=Hs.301667 Homo sapiens mRNA; cDNA DKFZp566I043 (from clon	AF037219	1.47
mitochondrial elongation factor G	NM_024996	1.47
laminin, beta 1	M20206	1.47
hypothetical protein FLJ13215	NM_025004	1.47
carboxypeptidase A3 (mast cell)	NM_001870	1.47
heparan sulfate 6-O-sulfotransferase	BF512553	1.47
KIAA0867 protein	AF312918	1.47
guanylate binding protein 1, interferon-inducible, 67kD	BC002666	1.47
alanine-glyoxylate aminotransferase (oxalosis I; hyperoxaluria I; glycolicaciduria; serine-pyruvate aminotransferase)	NM_016236	1.47
protein tyrosine phosphatase, non-receptor type 22 (lymphoid)	NM_012411	1.47
collagen, type I, alpha 1	K01228	1.47
DKFZP564O823 protein	NM_015393	1.47
melatonin receptor 1B	NM_005959	1.47
KIAA0931 protein	AK001854	1.47

WW domain-containing oxidoreductase	NM_016373	1.47
potassium voltage-gated channel, shaker-related subfamily, member 1 (episodic ataxia with myokymia)	NM_000217	1.47
testis-specific kinase substrate	NM_021733	1.47
postmeiotic segregation increased 2-like 4	D38500	1.47
estrogen receptor 2 (ER beta)	AB006589	1.47
neuregulin 1	NM_004495	1.47
pleckstrin homology-like domain, family A, member 1	NM_007350	1.47
DnaJ (Hsp40) homolog, subfamily C, member 8	AA843895	1.47
KIAA0775 gene product	NM_014726	1.47
phospholipase D1, phosphatidylcholine-specific	U38545	1.47
Consensus includes gb:AL121975 /DEF=Human DNA sequence from clone RP3-422B11 on chromosome 6p11.2-12.3 Contains ESTs, STSs and GSSs. Contains part of the PRIM2A gene for tw	AL121975	1.47
hypothetical protein FLJ20047	NM_017639	1.47
retinoic acid receptor, alpha	BE383139	1.47
Consensus includes gb:AF005487.1 /DEF=Homo sapiens MHC class II antigen (DRB6) mRNA, HLA-DRB6*0201 allele, sequence. /FEA=mRNA /DB_XREF=gi:5915893 /UG=Hs.167385 Homo	AF005487	1.47
erythrocyte membrane protein band 4.1-like 2	NM_001431	1.47
calcium-sensing receptor (hypocalciuric hypercalcemia 1, severe neonatal hyperparathyroidism)	D50855	1.47
pellino homolog 1 (Drosophila)	NM_020651	1.47
DNA segment on chromosome 6 (unique) 2654 expressed sequence	NM_012135	1.47
caspase 10, apoptosis-related cysteine protease	NM_001230	1.47
epidermal growth factor receptor (erythroblastic leukemia viral (v-erb-b) oncogene homolog, avian)	K03193	1.47
eukaryotic translation initiation factor 1A	AL079283	1.47
ribosome binding protein 1 homolog 180kD (dog)	AF006751	1.47
fatty-acid-Coenzyme A ligase, long-chain 6	AF129166	1.47
hypothetical protein FLJ12439	NM_023077	1.47
peptide YY	NM_004160	1.47
copine VI (neuronal)	AL050397	1.47
discs, large (Drosophila) homolog-associated protein 2	AF119817	1.47
protein phosphatase 2 (formerly 2A), regulatory subunit A (PR 65), beta isoform	AW009884	1.47
somatostatin receptor-interacting protein	NM_016148	1.47
GATA binding protein 2	AL563460	1.47
hairy homolog (Drosophila)	BE973687	1.47
DNase II-like acid DNase	NM_021233	1.47
carbohydrate (chondroitin 6/keratan) sulfotransferase 3	AB017915	1.47
retinoic acid- and interferon-inducible protein (58kD)	NM_012420	1.47
CTAGE-1 protein	NM_022663	1.47
telomeric repeat binding factor (NIMA-interacting) 1	AI347136	1.46
Consensus includes gb:AF238870.1 /DEF=Homo sapiens clone GLSH-3 similar to gliadin mRNA sequence. /FEA=mRNA /DB_XREF=gi:8272577 /UG=Hs.275706 Homo sapiens clone GLSH-3	AF238870	1.46
doublecortex; lissencephaly, X-linked (doublecortin)	NM_000555	1.46
matrix metalloproteinase 13 (collagenase 3)	NM_002427	1.46
uridine monophosphate synthetase (orotate phosphoribosyl transferase and orotidine-5'-decarboxylase)	NM_000373	1.46
Consensus includes gb:AK024976.1 /DEF=Homo sapiens cDNA: FLJ21323 fis, clone COL02374. /FEA=mRNA /DB_XREF=gi:10437405 /UG=Hs.75914 Homo sapiens cDNA: FLJ21323 fis, clo	AK024976	1.46
Consensus includes gb:AL050032.1 /DEF=Homo sapiens mRNA; cDNA DKFZp566F1224 (from clone DKFZp566F1224). /FEA=mRNA /DB_XREF=gi:4884272 /UG=Hs.306307 Homo sapiens	AL050032	1.46
protein (peptidyl-prolyl cis/trans isomerase) NIMA-interacting, 4 (parvulin)	NM_006223	1.46
olfactory receptor, family 7, subfamily C, member 1	NM_017506	1.46
zinc-finger protein ZBRK1	NM_021632	1.46
cytochrome P450, subfamily IIA (phenobarbital-inducible), polypeptide 6	M33318	1.46
hypothetical protein FLJ20624	NM_017906	1.46
kalilikrein 13	NM_015596	1.46
RAB6C, member RAS oncogene family	AL136727	1.46
serine/threonine kinase 29	AJ006701	1.46
complement-c1q tumor necrosis factor-related protein; likely ortholog of mouse CORS26 (collagenous repeat-containing sequence of 26-kDa protein)	NM_030945	1.46
DKFZP564G092 protein	NM_015601	1.46
solute carrier family 17 (sodium-dependent inorganic phosphate cotransporter), member 7	NM_020309	1.46
Consensus includes gb:H49077 /FEA=EST /DB_XREF=gi:988918 /DB_XREF=est:yq18c05.s1 /CLONE=IMAGE:274160 /UG=Hs.159179 Homo sapiens mRNA sequence, IMAGE clone 446411	H49077	1.46
phosphodiesterase 4D, cAMP-specific (phosphodiesterase E3 dunce homolog, Drosophila)	R40917	1.46
PBX/knotted 1 homeobox 2	NM_022062	1.46
parvin, alpha	AK022316	1.46
bone morphogenetic protein 5	NM_021073	1.46
Down syndrome critical region gene 1-like 1	NM_005822	1.46
p300/CBP-associated factor	AV727449	1.46
cyclic nucleotide gated channel alpha 1	NM_000087	1.46
Consensus includes gb:AK024315.1 /DEF=Homo sapiens cDNA FLJ14253 fis, clone OVARC1001376. /FEA=mRNA /DB_XREF=gi:10436666 /UG=Hs.300694 Homo sapiens cDNA FLJ14253 fi	AK024315	1.46
hexosaminidase B (beta polypeptide)	NM_000521	1.46

RAN binding protein 7	AI741392	1.46
thyroglobulin	NM_003235	1.46
guanine nucleotide-releasing factor 2 (specific for crk proto-oncogene)	NM_005312	1.46
Consensus includes gb:BE742268 /FEA=EST /DB_XREF=gi:10156260 /DB_XREF=est:601575686F1 /CLONE=IMAGE:3836709 /UG=Hs.281706 sortilin 1	BE742268	1.46
rabphilin 3A-like (without C2 domains)	BC005153	1.46
potassium channel, subfamily K, member 15 (TASK-5)	NM_022358	1.46
Consensus includes gb:AL567940 /FEA=EST /DB_XREF=gi:12921802 /DB_XREF=est:AL567940 /CLONE=CS0DF036YK19 (3 prime) /UG=Hs.7967 ESTs	AL567940	1.46
hypothetical protein similar to preferentially expressed antigen of melanoma	AL022101	1.46
hypothetical protein DKFZp761K1824	NM_017597	1.46
solute carrier family 7, (cationic amino acid transporter, y+ system) member 11	NM_014331	1.46
hypothetical protein FLJ23129	NM_024763	1.46
chromosome 22 open reading frame 3	AA018187	1.46
hypothetical protein FLJ21276	NM_024633	1.46
EphA7	NM_004440	1.46
KIAA0480 gene product	AW299294	1.46
Consensus includes gb:BF222483 /FEA=EST /DB_XREF=gi:11129660 /DB_XREF=est:7p53a01.x1 /CLONE=IMAGE:3649417 /UG=Hs.274454 ESTs, Weakly similar to 2004399A chromosom	BF222483	1.46
KIAA0417 gene product	NM_025015	1.46
Consensus includes gb:S79281.1 /DEF=pancreatic ribonuclease human, mRNA Recombinant Partial, 491 nt. /FEA=mRNA /GEN=pancreatic ribonuclease, HP-RNase /PROD=pancreatic ribonu	S79281	1.46
Consensus includes gb:AL049218.1 /DEF=Homo sapiens mRNA; cDNA DKFZp564I1916 (from clone DKFZp564I1916). /FEA=mRNA /DB_XREF=gi:4499947 /UG=Hs.306291 Homo sapiens n	AL049218	1.46
hypothetical protein FLJ23311	NM_024680	1.46
methylmalonate-semialdehyde dehydrogenase	BG399778	1.46
transcription factor 7-like 2 (T-cell specific, HMG-box)	AI270770	1.46
Fc fragment of IgE, low affinity II, receptor for (CD23A)	NM_002002	1.46
Consensus includes gb:AL080161.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434N144 (from clone DKFZp434N144). /FEA=mRNA /DB_XREF=gi:5262623 /UG=Hs.210431 Homo sapiens mi	AL080161	1.46
gonadotropin-releasing hormone receptor	L03380	1.46
neural cell adhesion molecule 1	M22094	1.46
Consensus includes gb:BF109941 /FEA=EST /DB_XREF=gi:10939631 /DB_XREF=est:7172c05.x1 /CLONE=IMAGE:3527024 /UG=Hs.27183 Homo sapiens clone 23940 mRNA sequence	BF109941	1.46
Burkitt lymphoma receptor 1, GTP binding protein	NM_001716	1.46
Consensus includes gb:AW975117 /FEA=EST /DB_XREF=gi:8166322 /DB_XREF=est:EST387223 /UG=Hs.145501 ESTs, Weakly similar to ALU1_HUMAN ALU SUBFAMILY J SEQUENCE CONTAI	AW975117	1.46
phosphodiesterase 4D, cAMP-specific (phosphodiesterase E3 dunce homolog, Drosophila)	AF012074	1.46
Consensus includes gb:AA576497 /FEA=EST /DB_XREF=gi:2353997 /DB_XREF=est:nm76b07.s1 /CLONE=IMAGE:1074133 /UG=Hs.252497 ESTs	AA576497	1.46
hypothetical protein FLJ10761	NM_018208	1.46
hypothetical protein MGC11271	NM_024323	1.46
cell division cycle 2-like 5 (cholinesterase-related cell division controller)	BC001274	1.46
Consensus includes gb:AK021633.1 /DEF=Homo sapiens cDNA FLJ11571 fis, clone HEMBA1003328. /FEA=mRNA /DB_XREF=gi:10432853 /UG=Hs.306608 Homo sapiens cDNA FLJ11571 fi	AK021633	1.46
peptidyl-prolyl isomerase G (cyclophilin G)	AI638762	1.46
Consensus includes gb:AU146952 /FEA=EST /DB_XREF=gi:11008473 /DB_XREF=est:AU146952 /CLONE=HEMBB1001962 /UG=Hs.306624 Homo sapiens cDNA FLJ12046 fis, clone HEMBB	AU146952	1.46
solute carrier family 2 (facilitated glucose transporter), member 3	AA778684	1.46
X-ray repair complementing defective repair in Chinese hamster cells 2	NM_005431	1.46
G protein-coupled receptor kinase 6	AF040751	1.46
TGFB inducible early growth response	NM_005655	1.46
transient receptor potential channel 1	NM_003304	1.46
Consensus includes gb:BE552409 /FEA=EST /DB_XREF=gi:9794101 /DB_XREF=est:hy07f08.x1 /CLONE=IMAGE:3196647 /UG=Hs.280895 ESTs	BE552409	1.46
paired basic amino acid cleaving system 4	O28513	1.46
cytochrome P450, subfamily 46 (cholesterol 24-hydroxylase)	NM_006668	1.46
hypothetical protein MGC2718	BE503090	1.46
fucosyltransferase 6 (alpha (1,3) fucosyltransferase)	U27335	1.46
CGI-143 protein	NM_016074	1.46
transmembrane 4 superfamily member 1	AI346835	1.46
semanogelin I	NM_003007	1.46
chloride channel, calcium activated, family member 2	NM_006536	1.46
matrix metalloprotease 27	NM_022122	1.46
secreted protein, acidic, cysteine-rich (osteonectin)	NM_003118	1.46
Consensus includes gb:AW972359 /FEA=EST /DB_XREF=gi:8162205 /DB_XREF=est:EST384450 /UG=Hs.293334 ESTs	AW972359	1.46
ephrin-B2	U16797	1.46
homeo box B2	NM_002145	1.46
hypothetical protein FLJ10232	NM_018033	1.46
small inducible cytokine subfamily A (Cys-Cys), member 19	U88321	1.46
CATR tumorigenicity conversion 1	U25433	1.46
hypothetical protein PRO1617	NM_018587	1.46
fyn-related kinase	NM_002031	1.46
UDP-Gal:betaGlcNAc beta 1,4- galactosyltransferase, polypeptide 4	BF223021	1.46
synuclein, alpha (non A4 component of amyloid precursor)	NM_000345	1.46

centaurin, gamma 2	NM_014914	1.46
glutathione S-transferase M4	M99422	1.46
NPHS2 gene (podocin)	NM_014625	1.46
glucocorticoid modulatory element binding protein 2	AL133646	1.46
nuclear receptor coactivator 2	AW027474	1.46
erythropoietin	AF053356	1.46
par-3 partitioning defective 3 homolog (C. elegans)	AF196185	1.46
hippocalcin-like protein 4	BE550384	1.46
S100 calcium binding protein A3	NM_002960	1.46
glial cells missing homolog a (Drosophila)	AB026493	1.46
amyloid beta (A4) precursor protein-binding, family B, member 1 (Fe65)	NM_001164	1.46
protocadherin 22	AJ276803	1.46
DNA (cytosine-5-)-methyltransferase 3 beta	NM_006892	1.46
Consensus includes gb:BF032808 /FEA=EST /DB_XREF=gi:10740520 /DB_XREF=est:601455629F1 /CLONE=IMAGE:3859303 /UG=Hs.266784 ESTs	BF032808	1.46
cytochrome P450, subfamily IIC (mephenytoin 4-hydroxylase), polypeptide 8	NM_000770	1.46
hypothetical protein	NM_015370	1.46
hypothetical protein FLJ10260	NM_018042	1.46
Consensus includes gb:AL137475.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434F0723 (from clone DKFZp434F0723). /FEA=mRNA /DB_XREF=gi:6808086 /UG=Hs.274578 Homo sapiens	AL137475	1.46
ribosomal protein S6 kinase, 90kD, polypeptide 5	NM_004755	1.46
RAS guanyl releasing protein 2 (calcium and DAG-regulated)	AI688812	1.46
VAMP (vesicle-associated membrane protein)-associated protein B and C	NM_004738	1.46
sialyltransferase 8C (alpha2,3&alpha1,4GlcNAcalpha 2,8-sialyltransferase)	NM_015879	1.46
hippocalcin	BC001777	1.46
chromosome 11 open reading frame 25	AI300461	1.46
endothelin 3	NM_000114	1.46
hypothetical protein FLJ20126	NM_017677	1.46
tumor necrosis factor receptor superfamily, member 12 (translocating chain-association membrane protein)	U94506	1.46
calcium channel, voltage-dependent, gamma subunit 5	NM_014404	1.46
claudin 1	NM_021101	1.46
CD44 antigen (homing function and Indian blood group system)	AI493245	1.46
hypothetical protein	NM_016619	1.46
leukocyte immunoglobulin-like receptor, subfamily A (with TM domain), member 1	NM_006863	1.46
Consensus includes gb:AK025206.1 /DEF=Homo sapiens cDNA: FLJ21553 fis, clone COL06329. /FEA=mRNA /DB_XREF=gi:10437671 /UG=Hs.306786 Homo sapiens cDNA: FLJ21553 fis, cl	AK025206	1.46
sialyltransferase 8C (alpha2,3&alpha1,4GlcNAcalpha 2,8-sialyltransferase)	NM_015879	1.46
synaptonemal complex protein 2	NM_014258	1.46
DKFZP434D1335 protein	AW593859	1.46
gap junction protein, beta 5 (connexin 31.1)	NM_005268	1.46
arginase, liver	NM_000045	1.46
Consensus includes gb:AL050391.1 /DEF=Homo sapiens mRNA; cDNA DKFZp586A181 (from clone DKFZp586A181); partial cds. /FEA=mRNA /GEN=DKFZp586A181 /PROD=hypothetical pr	AL050391	1.46
M27830 Human 28S ribosomal RNA gene, complete cds (_5, _M, _3 represent transcript regions 5 prime, Middle, and 3 prime respectively)	M27830	1.46
zinc finger protein 286	NM_020652	1.46
6 antigen 3	NM_001473	1.46
polyadenylate binding protein-interacting protein 1	NM_006451	1.46
hypothetical protein FLJ11383	NM_024938	1.46
synuclein, alpha (non A4 component of amyloid precursor)	BG260394	1.46
tumor necrosis factor, alpha-induced protein 6	AW188198	1.46
cysteine and glycine-rich protein 2	U46006	1.46
phosphatidylinositol glycan, class C	NM_002642	1.46
chemokine-like receptor 1	NM_004072	1.46
Consensus includes gb:AV652031 /FEA=EST /DB_XREF=gi:9873045 /DB_XREF=est:AV652031 /CLONE=GLCCWA12 /UG=Hs.282344 Homo sapiens cDNA FLJ13387 fis, clone PLACE1001131	AV652031	1.46
protocadherin gamma subfamily A, 1	AF152507	1.46
tripartite motif-containing 5	AF220028	1.46
nuclear receptor interacting protein 1	AI824012	1.46
mannan-binding lectin serine protease 2	NM_006610	1.46
proteoglycan 4, (megakaryocyte stimulating factor, articular superficial zone protein, camptodactyly, arthropathy, coxa vara, pericarditis syndrome)	NM_005807	1.45
plakophilin 2	AA888057	1.45
Consensus includes gb:AL530331 /FEA=EST /DB_XREF=gi:12793824 /DB_XREF=est:AL530331 /CLONE=CS0DD007YM06 (3 prime) /UG=Hs.111243 MADS box transcription enhancer facto	AL530331	1.45
thymidine kinase 2, mitochondrial	BE895437	1.45
Consensus includes gb:AI005205 /DEF=Homo sapiens 5HT3 gene for serotonin 3 receptor /FEA=mRNA_2 /DB_XREF=gi:7019744 /UG=Hs.2142 5-hydroxytryptamine (serotonin) receptor	AI005205	1.45
phospholipase A2, group IID	AL158172	1.45
Consensus includes gb:AL390157.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434D179 (from clone DKFZp434D179). /FEA=mRNA /DB_XREF=gi:9368847 /UG=Hs.306520 Homo sapiens mI	AL390157	1.45
HT001 protein	NM_014065	1.45
BCL2-associated athanogene 4	NM_004874	1.45

gb:NM_019841.1 /DEF=Homo sapiens epithelial calcium channel 1 (ECAC1), mRNA. /FEA=mRNA /GEN=ECAC1 /PROD=epithelial calcium channel 1 /DB_XREF=gi:9789940 /UG=Hs.283369	NM_019841	1.45
G protein-coupled receptor 56	AL554008	1.45
T cell receptor delta locus	AW007751	1.45
transmembrane protein 5	BF224146	1.45
endothelial differentiation, lysophosphatidic acid G-protein-coupled receptor, 2	BF055366	1.45
UDP-Gal:betaGlcNAc beta 1,4- galactosyltransferase, polypeptide 2	BC002431	1.45
Consensus includes gb:AU148957 /FEA=EST /DB_XREF=gi:11010478 /DB_XREF=est:AU148957 /CLONE=NT2RM4001313 /UG=Hs.306638 Homo sapiens cDNA FLJ12591 fis, clone NT2RM	AU148957	1.45
sarcospan (Kras oncogene-associated gene)	AL136756	1.45
heparan sulfate (glucosamine) 3-O-sulfotransferase 3A1	NM_006042	1.45
vinexin beta (SH3-containing adaptor molecule-1)	NM_005775	1.45
likely ortholog of mouse SRY-box containing gene 17	NM_022454	1.45
phospholipase A2, group VI (cytosolic, calcium-independent)	NM_003560	1.45
hypothetical protein PRO2792	AF119898	1.45
apolipoprotein C-III	NM_000040	1.45
myosin binding protein H	NM_004997	1.45
hypothetical protein FLJ20094	NM_017665	1.45
transducin beta-like 1 protein	AF332222	1.45
Prader-Willi/Angelman syndrome-5	AU118874	1.45
hypothetical protein TCBAP0758	AF283769	1.45
host cell factor C1 (VP16-accessory protein)	AA703045	1.45
glycoprotein hormones, alpha polypeptide	NM_000735	1.45
Consensus includes gb:AL080068.1 /DEF=Homo sapiens mRNA; cDNA DKFZp564J062 (from clone DKFZp564J062). /FEA=mRNA /DB_XREF=gi:5262475 /UG=Hs.272534 Homo sapiens mRI	AL080068	1.45
myotubularin related protein 3	AF233437	1.45
endothelin converting enzyme-like 1	NM_004826	1.45
myeloid/lymphoid or mixed-lineage leukemia (trithorax homolog, Drosophila); translocated to, 4	NM_005936	1.45
cartilage associated protein	NM_006371	1.45
serologically defined breast cancer antigen 84	AF308298	1.45
putative zinc finger protein from EUROIMAGE 566589	BC006148	1.45
H3 histone, family 3A	AI955655	1.45
guanylate cyclase 1, soluble, beta 3	W93728	1.45
LIM domain binding 3	AI803302	1.45
KIAA0417 gene product	AB007877	1.45
hypothetical protein FLJ11535	NM_024888	1.45
mucin 3B	AB038783	1.45
eukaryotic translation elongation factor 1 delta (guanine nucleotide exchange protein)	AL080315	1.45
growth hormone 1	NM_022559	1.45
metallothionein 1E (functional)	BF217861	1.45
cytochrome P450, subfamily IIIA, polypeptide 7	NM_000765	1.45
Consensus includes gb:AK024173.1 /DEF=Homo sapiens cDNA FLJ14111 fis, clone MAMMA1001630. /FEA=mRNA /DB_XREF=gi:10436489 /UG=Hs.288567 Homo sapiens cDNA FLJ14111	AK024173	1.45
Consensus includes gb:AK024129.1 /DEF=Homo sapiens cDNA FLJ14067 fis, clone HEMBB1001315. /FEA=mRNA /DB_XREF=gi:10436434 /UG=Hs.287619 Homo sapiens cDNA FLJ14067 f	AK024129	1.45
SON DNA binding protein	AI936458	1.45
putative GTP-binding protein similar to RAY/RAB1C	AL037167	1.45
hypothetical protein FLJ21463	AK025116	1.45
growth hormone receptor	NM_000163	1.45
KIAA0528 gene product	AB011100	1.45
Consensus includes gb:AK026273.1 /DEF=Homo sapiens cDNA: FLJ22620 fis, clone HSI05629. /FEA=mRNA /DB_XREF=gi:10439077 /UG=Hs.306834 Homo sapiens cDNA: FLJ22620 fis, cl	AK026273	1.45
carbonic anhydrase IV	NM_000717	1.45
Consensus includes gb:AI017119 /FEA=EST /DB_XREF=gi:3231455 /DB_XREF=est:ov34a05.x1 /CLONE=IMAGE:1639184 /UG=Hs.144339 Human DNA sequence from clone RP1-257A15 on	AI017119	1.45
hypothetical protein MGC2656	NM_024509	1.45
synovial sarcoma, X breakpoint 3	BC005904	1.45
28kD interferon responsive protein	NM_022147	1.45
gb:Z25427.1 /DEF=H.sapiens protein-serinethreonine kinase gene, complete CDS. /FEA=mRNA /PROD=protein-serinethreonine kinase /DB_XREF=gi:405732 /FL=gb:Z25427.1	Z25427	1.45
glutamate receptor, ionotropic, kainate 1	NM_000830	1.45
Consensus includes gb:AL049430.1 /DEF=Homo sapiens mRNA; cDNA DKFZp586H201 (from clone DKFZp586H201). /FEA=mRNA /DB_XREF=gi:4500210 /UG=Hs.302061 Homo sapiens mI	AL049430	1.45
PRO0457 protein	NM_021651	1.45
f-box and WD-40 domain protein 18	AB033281	1.45
Consensus includes gb:BE378994 /FEA=EST /DB_XREF=gi:9324359 /DB_XREF=est:601236247F1 /CLONE=IMAGE:3608626 /UG=Hs.271717 ESTs, Weakly similar to ALU1_HUMAN ALU SUI	BE378994	1.45
calponin 3, acidic	NM_001839	1.45
maltase-glucoamylase (alpha-glucosidase)	NM_004668	1.45
DKFZP434F122 protein	AL080200	1.45
transporter 2, ATP-binding cassette, sub-family B (MDR/TAP)	M74447	1.45
sialic acid binding Ig-like lectin 6	NM_001245	1.45
protein tyrosine phosphatase, receptor type, N polypeptide 2	AF007555	1.45

Consensus includes gb:AU158443 /FEA=EST /DB_XREF=gi:11019964 /DB_XREF=est:AU158443 /CLONE=PLACE2000398 /UG=Hs.287592 Homo sapiens cDNA FLJ13713 fis, clone PLACE2(	AU158443	1.45
E74-like factor 3 (ets domain transcription factor, epithelial-specific)	AF017307	1.45
H2B histone family, member Q	NM_003528	1.45
KIAA1751 protein	AB051538	1.45
Consensus includes gb:BF348061 /FEA=EST /DB_XREF=gi:11295656 /DB_XREF=est:602022610F1 /CLONE=IMAGE:4157907 /UG=Hs.167988 neural cell adhesion molecule 1	BF348061	1.45
v-kit Hardy-Zuckerman 4 feline sarcoma viral oncogene homolog	NM_000222	1.45
hypothetical protein FLJ14803	BE619220	1.45
ATPase, Ca++ transporting, plasma membrane 2	X63575	1.45
DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 34	AA039439	1.45
M-phase phosphoprotein 10 (U3 small nucleolar ribonucleoprotein)	AL545921	1.45
putative DNA/chromatin binding motif	NM_006618	1.45
aldehyde oxidase 1	NM_001159	1.45
deleted in azoospermia 4	AF248483	1.45
Consensus includes gb:AI912086 /FEA=EST /DB_XREF=gi:5631941 /DB_XREF=est:wd84f08.x1 /CLONE=IMAGE:2338311 /UG=Hs.194750 Human DNA sequence from clone 1170K4 on chr	AI912086	1.45
advillin	BC004134	1.45
chloride channel 1 , skeletal muscle (Thomsen disease, autosomal dominant)	NM_000083	1.45
H4 histone family, member J	NM_003545	1.45
nuclear factor (erythroid-derived 2), 45kD	L13974	1.45
hypothetical protein FLJ11585	AI082827	1.45
plasminogen activator, urokinase	K03226	1.45
KIAA0841 protein	AA845355	1.45
cullin 5	AF017061	1.45
VGF nerve growth factor inducible	NM_003378	1.45
latrophilin	NM_012302	1.45
glutamate decarboxylase 2 (pancreatic islets and brain, 65kD)	X69936	1.45
apolipoprotein (a) related gene C	U19518	1.45
pyruvate dehydrogenase (lipoamide) alpha 2	NM_005390	1.45
surfactant protein A binding protein	NM_005129	1.45
pregnancy-associated plasma protein A	AA148534	1.45
RAB33A, member RAS oncogene family	NM_004794	1.45
KIAA0644 gene product	NM_014817	1.45
KIAA0227 protein	AW235608	1.45
butyrophilin-like 3	AK025267	1.45
GATA binding protein 1 (globin transcription factor 1)	M30601	1.45
arginine-glutamic acid dipeptide (RE) repeats	AI920976	1.45
PRKC, apoptosis, WT1, regulator	BF732462	1.45
solute carrier family 1 (neuronal/epithelial high affinity glutamate transporter, system Xag), member 1	NM_004170	1.45
runt-related transcription factor 1 (acute myeloid leukemia 1; aml1 oncogene)	D89788	1.45
triple functional domain (PTPRF interacting)	AL161955	1.45
neurochondrin	AB018740	1.45
spondin 1, (f-spondin) extracellular matrix protein	AB051390	1.45
like-glycosyltransferase	NM_004737	1.45
chromosome 2 open reading frame 6	AK023321	1.45
hydroxy-delta-5-steroid dehydrogenase, 3 beta- and steroid delta-isomerase 1	NM_000862	1.45
hypothetical protein	NM_019040	1.45
Consensus includes gb:AW814026 /FEA=EST /DB_XREF=gi:7907020 /DB_XREF=est:RC6-ST0198-071199-021-602 /UG=Hs.273621 Homo sapiens cDNA: FLJ21350 fis, clone COL02751	AW814026	1.45
Consensus includes gb:BF739971 /FEA=EST /DB_XREF=gi:12066647 /DB_XREF=est:7o41f07.x1 /CLONE=IMAGE:3576732 /UG=Hs.98416 ESTs	BF739971	1.45
gb:L14455.1 /DEF=Human Ig rearranged mu-chain gene V-N-D-N-J-reion, complete cds. /FEA=mRNA /DB_XREF=gi:398496 /FL=gb:L14455.1	L14455	1.45
N-deacetylase/N-sulfotransferase (heparan glucosaminyl) 2	NM_003635	1.45
BAI1-associated protein 3	BE328802	1.45
paraoxonase 2	AF001602	1.45
KIAA1922 protein	AK026910	1.45
keratin, hair, acidic, 4	NM_021013	1.45
taste receptor, type 2, member 4	NM_016944	1.45
chromosome 11 open reading frame 8	BE645386	1.45
transmembrane protease, serine 2	AF270487	1.45
lipopolysaccharide binding protein	M35533	1.45
Consensus includes gb:AK023816.1 /DEF=Homo sapiens cDNA FLJ13754 fis, clone PLACE3000362. /FEA=mRNA /DB_XREF=gi:10435865 /UG=Hs.136233 Homo sapiens cDNA FLJ13754 fi	AK023816	1.45
G protein-coupled receptor 17	NM_005291	1.45
Consensus includes gb:AK000993.1 /DEF=Homo sapiens cDNA FLJ10131 fis, clone HEMBA1003041. /FEA=mRNA /DB_XREF=gi:7021996 /UG=Hs.274128 Homo sapiens cDNA FLJ10131 fis	AK000993	1.45
gb:NM_005179.1 /DEF=Homo sapiens avian lymphoma virus-derived transforming sequence (BLYM), mRNA. /FEA=CDS /GEN=BLYM /PROD=avian lymphoma virus-derived transformingseq	NM_005179	1.45
carboxylesterase-related protein	NM_016280	1.45
hypothetical protein DKFZp564A176	AL136663	1.45

alkaline phosphatase, placental (Regan isozyme)	NM_001632	1.45
matrilin 1, cartilage matrix protein	M55683	1.45
hypothetical protein DKFZp761K1423	NM_018422	1.45
gephyrin	NM_020806	1.45
leptin receptor	U66495	1.45
Consensus includes gb:AA558583 /FEA=EST /DB_XREF=gi:2329060 /DB_XREF=est:nk38h06.s1 /CLONE=IMAGE:1015835 /UG=Hs.288826 Homo sapiens cDNA FLJ12263 fis, clone MAMMA	AA558583	1.45
L38424 B subtilis dapB, jojF, joj6 genes corresponding to nucleotides 1358-3197 of L38424 (-5, -M, -3 represent transcript regions 5 prime, Middle, and 3 prime respectively)	L38424	1.45
regulator of G-protein signalling 11	AB016929	1.45
Consensus includes gb:AA070330 /FEA=EST /DB_XREF=gi:1577689 /DB_XREF=est:zm68e06.r1 /CLONE=IMAGE:530818 /UG=Hs.283729 ESTs	AA070330	1.45
hypothetical protein FLJ10945	NM_018280	1.45
hypothetical protein FLJ13511	AK023573	1.45
HBS1-like (S. cerevisiae)	AK024258	1.45
Consensus includes gb:AK024568.1 /DEF=Homo sapiens cDNA: FLJ20915 fis, clone ADSE00692. /FEA=mRNA /DB_XREF=gi:10436878 /UG=Hs.306691 Homo sapiens cDNA: FLJ20915 fis,	AK024568	1.44
brain abundant, membrane attached signal protein 1	NM_006317	1.44
midline 2	NM_012216	1.44
carcinoembryonic antigen-related cell adhesion molecule 7	NM_006890	1.44
KIAA1128 protein	AI478879	1.44
a disintegrin-like and metalloprotease (reprolysin type) with thrombospondin type 1 motif, 8	NM_007037	1.44
olfactory receptor, family 1, subfamily D, member 2	NM_002548	1.44
KIAA1644 protein	BF516072	1.44
regulator of G-protein signalling 13	AF030107	1.44
GAS2-related on chromosome 22	Y07846	1.44
arylsulfatase A	NM_000487	1.44
GTP binding protein 1	NM_004286	1.44
hypothetical protein from EUROIMAGE 511235	NM_020147	1.44
dedicator of cyto-kinesis 1	NM_001380	1.44
proteasome (prosome, macropain) subunit, beta type, 1	W86293	1.44
phospholipase A2 receptor 1, 180kD	U17033	1.44
KRAB box containing C2H2 type zinc finger bA526D8.4	BE567977	1.44
Huntingtin interacting protein B	NM_014159	1.44
CD36 antigen, gamma polypeptide (TiT3 complex)	NM_000073	1.44
paired box gene 3 (Waardenburg syndrome 1)	U02309	1.44
hypothetical protein FLJ13031	NM_024688	1.44
protein phosphatase 3 (formerly 2B), catalytic subunit, gamma isoform (calcineurin A gamma)	BC004864	1.44
protein with polyglutamine repeat; calcium (ca2+) homeostasis endoplasmic reticulum protein	AI372979	1.44
keratin associated protein 4.7	AI406939	1.44
origin recognition complex, subunit 6 (yeast homolog)-like	AF336878	1.44
KIAA0506 protein	AB007975	1.44
sialophorin (gpL115, leukosialin, CD43)	J04168	1.44
endothelial differentiation, lysophosphatidic acid G-protein-coupled receptor, 4	AF011466	1.44
spindlin-like	NM_019003	1.44
Consensus includes gb:BE886287 /FEA=EST /DB_XREF=gi:10340242 /DB_XREF=est:601509871F1 /CLONE=IMAGE:3911608 /UG=Hs.249219 Homo sapiens chromosome 19, overlapping	BE886287	1.44
dual specificity phosphatase 9	NM_001395	1.44
duodenal cytochrome b	NM_024843	1.44
hypothetical protein FLJ11756	NM_024606	1.44
glucocorticoid receptor DNA binding factor 1	NM_004491	1.44
Consensus includes gb:AV721430 /FEA=EST /DB_XREF=gi:10818582 /DB_XREF=est:AV721430 /CLONE=HTBBIF09 /UG=Hs.173638 transcription factor 7-like 2 (T-cell specific, HMG-box)	AV721430	1.44
hypothetical protein FLJ13381	NM_025068	1.44
v-ets erythroblastosis virus E26 oncogene homolog 2 (avian)	AL575509	1.44
cytoskeleton-associated protein 4	NM_006825	1.44
Consensus includes gb:AV647366 /FEA=EST /DB_XREF=gi:9868380 /DB_XREF=est:AV647366 /CLONE=GLCAXE09 /UG=Hs.282365 ESTs	AV647366	1.44
hypothetical protein FLJ20456	NM_017831	1.44
Consensus includes gb:AU159040 /FEA=EST /DB_XREF=gi:11020561 /DB_XREF=est:AU159040 /CLONE=THYRO1000676 /UG=Hs.296745 Homo sapiens cDNA FLJ13833 fis, clone THYRO	AU159040	1.44
HFSE-1 protein	AF072164	1.44
Consensus includes gb:AL353942.1 /DEF=Homo sapiens mRNA; cDNA DKFZp761L23121 (from clone DKFZp761L23121). /FEA=mRNA /DB_XREF=gi:7669981 /UG=Hs.306504 Homo sapie	AL353942	1.44
Consensus includes gb:AL050388.1 /DEF=Homo sapiens mRNA; cDNA DKFZp564M2422 (from clone DKFZp564M2422); partial cds. /FEA=mRNA /GEN=DKFZp564M2422 /PROD=hypothetic	AL050388	1.44
tripartite motif-containing 2	BC005016	1.44
zinc finger protein 239	NM_005674	1.44
hypothetical protein FLJ11362	NM_021946	1.44
hypothetical protein FLJ23558	NM_025095	1.44
tumor necrosis factor (ligand) superfamily, member 4 (tax-transcriptionally activated glycoprotein 1, 34kD)	NM_003326	1.44
SEC24 related gene family, member A (S. cerevisiae)	AI131244	1.44
UL16 binding protein 1	NM_025218	1.44

KIAA0892 protein	AC003030	1.44
POU domain, class 4, transcription factor 1	L20433	1.44
solute carrier family 6 (neurotransmitter transporter, creatine), member 8	NM_005629	1.44
KIAA0773 gene product	NM_014690	1.44
hypothetical protein	BE779865	1.44
myosin, heavy polypeptide 3, skeletal muscle, embryonic	NM_002470	1.44
Consensus includes gb:AL137428.1 /DEF=Homo sapiens mRNA; cDNA DKFZp761N1323 (from clone DKFZp761N1323). /FEA=mRNA /DB_XREF=gi:6807985 /UG=Hs.306459 Homo sapiens	AL137428	1.44
KIAA0608 protein	N34407	1.44
Consensus includes gb:AA156240 /FEA=EST /DB_XREF=gi:1727858 /DB_XREF=est:zl50c12.s1 /CLONE=IMAGE:505366 /UG=Hs.288660 Homo sapiens cDNA: FLJ22182 fis, clone HRC0095	AA156240	1.44
solute carrier family 17 (sodium phosphate), member 3	NM_006632	1.44
growth hormone 1	NM_022562	1.44
dentatorubral-pallidoluysian atrophy (atrophin-1)	D31840	1.44
adenylate kinase 3	NM_013410	1.44
stratifin	X57348	1.44
hypothetical protein FLJ13479	NM_024706	1.44
serine (or cysteine) proteinase inhibitor, clade E (nexin, plasminogen activator inhibitor type 1), member 1	NM_000602	1.44
hypothetical protein FLJ20202	NM_017709	1.44
BTG family, member 3	AL049332	1.44
prostaglandin E receptor 2 (subtype EP2), 53kD	NM_000956	1.44
CD24 antigen (small cell lung carcinoma cluster 4 antigen)	AA761181	1.44
ribonuclease, RNase A family, 4	NM_002937	1.44
stathmin 1/oncoprotein 18	NM_005563	1.44
relaxin 1 (H1)	BC005956	1.44
hBOAT for potent brain type organic ion transporter	NM_016609	1.44
homeo box A10	BF792917	1.44
RAP1, GTPase activating protein 1	AB007943	1.44
KIAA0871 protein	AA861784	1.44
v-ral simian leukemia viral oncogene homolog A (ras related)	NM_005402	1.44
phosphodiesterase 8B	AK023913	1.44
T-box 6	AK022330	1.44
a disintegrin and metalloproteinase domain 5	AI132821	1.44
KIAA1919 protein	AK000168	1.44
integrin, alpha L (antigen CD11A (p180), lymphocyte function-associated antigen 1; alpha polypeptide)	AC002310	1.44
cyclin-dependent kinase inhibitor 1C (p57, Kip2)	N33167	1.44
small 1 homolog, zinc finger protein (Drosophila)	NM_005985	1.44
KIAA0856 protein	AB020663	1.44
palladin	NM_016081	1.44
hypothetical protein FLJ11583	NM_024663	1.44
leukocyte immunoglobulin-like receptor, subfamily B (with TM and ITIM domains), member 1	NM_006669	1.44
ATP-binding cassette, sub-family C (CFTR/MRP), member 8	L78207	1.44
glutamate receptor, ionotropic, N-methyl D-aspartate 2B	U90278	1.44
Consensus includes gb:U81961 /DEF=Homo sapiens amiloride-sensitive epithelial sodium channel alpha subunit alpha ENaC-1 and alpha ENaC-2 (SCNN1A) genes, alternatively spliced, parti	U81961	1.44
cytochrome P450, subfamily IIB (phenobarbital-inducible), polypeptide 6	NM_000767	1.44
Consensus includes gb:AF090910.1 /DEF=Homo sapiens clone HQ0262. /FEA=mRNA /DB_XREF=gi:6690195 /UG=Hs.236760 Homo sapiens clone HQ0262	AF090910	1.44
vacuolar protein sorting 11 (yeast)	NM_021729	1.44
PSR2b protein	NM_016559	1.44
leukocyte immunoglobulin-like receptor, subfamily B (with TM and ITIM domains), member 4	U82979	1.44
serum/glucocorticoid regulated kinase 2	Z98752	1.44
RAB11B, member RAS oncogene family	AL575337	1.44
Gli-interacting zinc finger protein	AA054734	1.44
inositol 1,4,5-triphosphate receptor, type 3	BF001241	1.44
Consensus includes gb:BF592058 /FEA=EST /DB_XREF=gi:11684382 /DB_XREF=est:7n98g09.x1 /CLONE=IMAGE:3572920 /UG=Hs.86405 Homo sapiens mRNA; cDNA DKFZp564P056 (fro	BF592058	1.44
synaptojanin 2	AA191573	1.44
insulin-like growth factor 1 (somatomedin C)	AI972496	1.44
sex comb on midleg homolog 1	NM_012236	1.44
sema domain, immunoglobulin domain (Ig), and GPI membrane anchor, (semaphorin) 7A	AF071542	1.44
tachykinin receptor 3	NM_001059	1.44
hypothetical protein FLJ22595	NM_025047	1.44
adenovirus 5 E1A binding protein	NM_006624	1.44
ATP-binding cassette, sub-family A (ABC1), member 2	AF327657	1.44
serine protease inhibitor, Kunitz type, 3	AW118166	1.44
UDP-glucuronic acid/UDP-N-acetylgalactosamine dual transporter	AI769637	1.44
neurofibromin 1 (neurofibromatosis, von Recklinghausen disease, Watson disease)	D12625	1.44

CD68 antigen	NM_001251	1.44
keratin 4	X07695	1.44
zinc finger protein 230	NM_006300	1.44
cisplatin resistance associated	NM_006697	1.44
popeye protein 2	NM_022135	1.44
I-mfa domain-containing protein	AF054589	1.44
olfactory receptor, family 1, subfamily E, member 2	NM_003554	1.44
hypothetical protein FLJ13840	NM_024746	1.44
ELAV (embryonic lethal, abnormal vision, Drosophila)-like 2 (Hu antigen B)	NM_004432	1.44
hypothetical protein MGC2827	NM_023940	1.44
hypothetical protein FLJ14351	NM_024732	1.44
transmembrane 4 superfamily member 6	AF053453	1.44
Notch homolog 3 (Drosophila)	NM_000435	1.44
Consensus includes gb:AK022102.1 /DEF=Homo sapiens cDNA FLJ12040 fis, clone HEMBB1001944. /FEA=mRNA /DB_XREF=gi:10433423 /UG=Hs.296687 Homo sapiens cDNA FLJ12040 f	AK022102	1.44
allograft inflammatory factor 1	NM_004847	1.44
insulin receptor substrate 1	NM_005544	1.44
Consensus includes gb:BE044440 /FEA=EST /DB_XREF=gi:8361493 /DB_XREF=est:ho45e07.s1 /CLONE=IMAGE:3040356 /UG=Hs.251278 KIAA1201 protein	BE044440	1.44
pervin, beta	N73272	1.44
neurofibromin 1 (neurofibromatosis, von Recklinghausen disease, Watson disease)	M60915	1.44
phospholipase A2, group IB (pancreas)	NM_000928	1.44
Cas-Br-M (murine) ectropic retroviral transforming sequence b	U26710	1.44
calcium binding protein 1 (calbrain)	NM_004276	1.44
Consensus includes gb:AF005081.1 /DEF=Homo sapiens skin-specific protein (sp32) mRNA, partial cds. /FEA=mRNA /GEN=sp32 /PROD=skin-specific protein /DB_XREF=gi:2589189 /UG=	AF005081	1.44
growth arrest-specific 7	BC001152	1.44
KIAA0472 protein	AF068286	1.44
dynein, cytoplasmic, intermediate polypeptide 1	NM_004411	1.44
PRO1900 protein	NM_016344	1.44
EGF-containing fibulin-like extracellular matrix protein 2	AB030655	1.44
monoamine oxidase B	NM_000898	1.44
Consensus includes gb:AU147182 /FEA=EST /DB_XREF=gi:11008703 /DB_XREF=est:AU147182 /CLONE=HEMBB1002623 /UG=Hs.301197 Homo sapiens cDNA FLJ12098 fis, clone HEMBB	AU147182	1.44
Consensus includes gb:AL442084.1 /DEF=Homo sapiens mRNA; cDNA DKFZp547G044 (from clone DKFZp547G044). /FEA=mRNA /DB_XREF=gi:10241763 /UG=Hs.306924 Homo sapiens n	AL442084	1.44
hypothetical protein FLJ11286	AI862559	1.44
guanylate kinase 1	AW182892	1.44
hypothetical protein FLJ13231	NM_023073	1.44
growth differentiation factor 11	AF028333	1.44
Consensus includes gb:X07868 /DEF=Human DNA for insulin-like growth factor II (IGF-2); exon 7 and additional ORF /FEA=mRNA_1 /DB_XREF=gi:32998 /UG=Hs.251664 insulin-like growt	X07868	1.44
frizzled homolog 4 (Drosophila)	NM_012193	1.44
KIAA0599 protein	AU147160	1.44
renal tumor antigen	NM_014226	1.44
KIAA1048 protein	BC002695	1.44
guanylate binding protein 2, interferon-inducible	NM_004120	1.44
SAM domain, SH3 domain and nuclear localisation signals, 1	NM_022136	1.44
cathepsin B	NM_001908	1.44
Consensus includes gb:BE783279 /FEA=EST /DB_XREF=gi:10204477 /DB_XREF=est:601470835F1 /CLONE=IMAGE:3873801 /UG=Hs.33862 ESTs	BE783279	1.44
ectodysplasin A2 isoform receptor	NM_021783	1.44
KIAA0853 protein	BE895685	1.44
potassium voltage-gated channel, KQT-like subfamily, member 2	AF110020	1.44
tissue inhibitor of metalloproteinase 3 (Sorsby fundus dystrophy, pseudoinflammatory)	NM_000362	1.44
hypothetical protein FKSG28	W28828	1.43
Consensus includes gb:N30878 /FEA=EST /DB_XREF=gi:1149398 /DB_XREF=est:yw76b05.s1 /CLONE=IMAGE:258129 /UG=Hs.82276 trichohyalin	N30878	1.43
solute carrier family 21 (organic anion transporter), member 14	NM_017435	1.43
CD36 antigen (collagen type I receptor, thrombospondin receptor)-like 2 (lysosomal integral membrane protein II)	NM_005506	1.43
slit homolog 3 (Drosophila)	AB011538	1.43
v-akt murine thymoma viral oncogene homolog 1	NM_005163	1.43
platelet/endothelial cell adhesion molecule (CD31 antigen)	AA702701	1.43
epithelial membrane protein 1	BF445047	1.43
Consensus includes gb:AL021327 /DEF=Human DNA sequence from PAC 12409 on chromosome 6q21. Contains DNAJ2 (HDJ1) like pseudogene, ESTs, STSs and GSSs /FEA=CDS /DB_XREF=	AL021327	1.43
Consensus includes gb:AK024457.1 /DEF=Homo sapiens mRNA for FLJ00049 protein, partial cds. /FEA=mRNA /GEN=FLJ00049 /PROD=FLJ00049 protein /DB_XREF=gi:10440427 /UG=Hs.	AK024457	1.43
tubulin, gamma 2	NM_016437	1.43
Consensus includes gb:S67777 /DEF=estrogen receptor (exon 6) human, tamoxifen-resistant breast tumor 17, Genomic Mutant, 187 nt /FEA=CDS_1 /DB_XREF=gi:459362 /UG=Hs.247976	S67777	1.43
secretagogin	NM_006998	1.43
X04603 B subtilis thrC, thrB genes corresponding to nucleotides 248-2229 of X04603 (-5, -M, -3 represent transcript regions 5 prime, Middle, and 3 prime respectively)	X04603	1.43
Consensus includes gb:AL034450 /DEF=Human DNA sequence from clone 115K14 on chromosome Xq22.3-23 Contains high mobility group protein 2a, ESTs, STS /FEA=mRNA /DB_XREF=gi	AL034450	1.43

transcription factor (p38 interacting protein)	AA889653	1.43
fibronectin 1	AI276395	1.43
bromodomain-containing 2	AA902767	1.43
ATPase, Ca++ transporting, plasma membrane 2	M97260	1.43
KIAA1077 protein	AI479175	1.43
Consensus includes gb:AV741657 /FEA=EST /DB_XREF=gi:10859238 /DB_XREF=est:AV741657 /CLONE=CBMALG01 /UG=Hs.154085 leucine zipper protein 1	AV741657	1.43
transient receptor potential channel 6	AI271068	1.43
HIV-1 inducer of short transcripts binding protein; lymphoma related factor	AW027070	1.43
gb:L06102.1 /DEF=Human IG processed VH-region mRNA, complete cds. /FEA=mRNA /GEN=IGH@ /PROD=immunoglobulin heavy chain V region /DB_XREF=gi:185528 /FL=gb:L06102.1	L06102	1.43
melanoma cell adhesion molecule	M28882	1.43
aldo-keto reductase family 1, member C4 (chlordecone reductase; 3-alpha hydroxysteroid dehydrogenase, type I; dihydrodiol dehydrogenase 4)	AB045829	1.43
hypothetical protein	U79275	1.43
tankyrase, TRF1-interacting ankyrin-related ADP-ribose polymerase	AF082559	1.43
contactin associated protein-like 2	NM_014141	1.43
chromosome X open reading frame 2	NM_001586	1.43
KIAA0276 protein	AA194584	1.43
villin 1	NM_007127	1.43
hydroxysteroid (11-beta) dehydrogenase 1	NM_005525	1.43
gb:NM_015977.1 /DEF=Homo sapiens Williams-Beuren syndrome chromosome region 14 (WBSCR14), mRNA. /FEA=mRNA /GEN=WBSCR14 /PROD=WBSCR14 protein /DB_XREF=gi:770574	NM_015977	1.43
hypothetical protein FLJ20004	NM_017616	1.43
single-minded homolog 2 (Drosophila)	NM_005069	1.43
hypothetical protein LOC57187	BE543527	1.43
protein phosphatase 3 (formerly 2B), regulatory subunit B (19kD), alpha isoform (calcineurin B, type I)	AL544951	1.43
NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 3 (12kD, B12)	NM_002491	1.43
tropomyosin 2 (beta)	AL566786	1.43
membrane protein, palmitoylated 2 (MAGUK p55 subfamily member 2)	NM_005374	1.43
Consensus includes gb:AK021640.1 /DEF=Homo sapiens cDNA FLJ11578 fis, clone HEMBA1003571. /FEA=mRNA /DB_XREF=gi:10432861 /UG=Hs.287427 Homo sapiens cDNA FLJ11578 f	AK021640	1.43
corticotropin releasing hormone receptor 1	AF180301	1.43
KIAA0509 protein	AB007978	1.43
PRO0246 protein	NM_014123	1.43
proliferation-associated 2G4, 38kD	BE501952	1.43
KIAA0555 gene product	AI962693	1.43
small inducible cytokine subfamily A (Cys-Cys), member 22	NM_002990	1.43
Consensus includes gb:D16471.1 /DEF=Human mRNA, Xq terminal portion. /FEA=mRNA /DB_XREF=gi:495099 /UG=Hs.121571 Human mRNA, Xq terminal portion	D16471	1.43
phosphatidic acid phosphatase type 2B	AV725664	1.43
hypothetical protein FLJ22390	NM_022746	1.43
homolog of rat orphan transporter v7-3	NM_018057	1.43
melanoma antigen, family A, 9	BC002351	1.43
hypothetical protein DKFZp547M136 similar to widely-interspaced zinc finger motifs	W67467	1.43
hypothetical protein FLJ20260	NM_017731	1.43
cyclin-dependent kinase 3	NM_001258	1.43
KIAA1750 protein	AI096375	1.43
Consensus includes gb:AI633741 /FEA=EST /DB_XREF=gi:4685071 /DB_XREF=est:tt28a05.x1 /CLONE=IMAGE:2242064 /UG=Hs.270364 ESTs, Moderately similar to S14458 laminin alpha	AI633741	1.43
myelin transcription factor 1-like	BF223003	1.43
Consensus includes gb:AK024457.1 /DEF=Homo sapiens mRNA for FLJ00049 protein, partial cds. /FEA=mRNA /GEN=FLJ00049 /PROD=FLJ00049 protein /DB_XREF=gi:10440427 /UG=Hs.	AK024457	1.43
protocadherin alpha 3	AF152481	1.43
E3 ubiquitin ligase SMURF1	AF199364	1.43
non-metastatic cells 5, protein expressed in (nucleoside-diphosphate kinase)	NM_003551	1.43
hypothetical protein	BC005004	1.43
ets variant gene 1	NM_004956	1.43
plexin C1	NM_005761	1.43
hypothetical protein FLJ23505	NM_024716	1.43
multiple PDZ domain protein	NM_003829	1.43
leucine-rich repeat-containing 3	NM_030891	1.43
sialidase 3 (membrane sialidase)	AK022450	1.43
RAN binding protein 8	NM_006390	1.43
apolipoprotein A-II	NM_001643	1.43
zinc finger protein 143 (clone pHZ-1)	NM_003442	1.43
H1 histone family, member 5	NM_005322	1.43
transition protein 2 (during histone to protamine replacement)	X63759	1.43
Rho guanine exchange factor (GEF) 15	AK023853	1.43
acetylserotonin O-methyltransferase-like	Y15521	1.43
runt-related transcription factor 2	NM_004348	1.43

sarcosine dehydrogenase	NM_007101	1.43
transcription factor 3 (E2A immunoglobulin enhancer binding factors E12/E47)	BG393795	1.43
gb:AF119892.1 /DEF=Homo sapiens PRO2710 mRNA, complete cds. /FEA=mRNA /PROD=PRO2710 /DB_XREF=gi:7770220 /UG=Hs.306572 Homo sapiens PRO2710 mRNA, complete cds /	AF119892	1.43
hypothetical protein FLJ23056	NM_024582	1.43
KIAA0409 protein	X72882	1.43
PRO0132 protein	NM_014116	1.43
Consensus includes gb:AI419423 /FEA=EST /DB_XREF=gi:4265354 /DB_XREF=est:tf28d09.s1 /CLONE=IMAGE:2097521 /UG=Hs.172780 ESTs	AI419423	1.43
hypothetical protein MGC3130	NM_024032	1.43
guanylate cyclase activator 1A (retina)	L36861	1.43
leucine zipper-EF-hand containing transmembrane protein 1	NM_012318	1.43
Consensus includes gb:W72053 /FEA=EST /DB_XREF=gi:1382323 /DB_XREF=est:zd66e04.s1 /CLONE=IMAGE:345630 /UG=Hs.110771 Homo sapiens cDNA: FLJ21904 fis, clone HEP03585	W72053	1.43
SH3 domain binding glutamic acid-rich protein	NM_007341	1.43
Sjogren syndrome antigen A1 (52kD, ribonucleoprotein autoantigen SS-A/Ro)	NM_003141	1.43
macrophage lectin 2 (calcium dependent)	NM_006344	1.43
paired box gene 8	AI948472	1.43
myosin IXB	NM_004145	1.43
chromosome 11 open reading frame 14	NM_020645	1.43
plasminogen activator, urokinase receptor	X74039	1.43
sulfotransferase family, cytosolic, 2A, dehydroepiandrosterone (DHEA) -preferring, member 1	U08024	1.43
phosphoinositide-3-kinase, class 2, gamma polypeptide	AJ000008	1.43
cholinergic receptor, nicotinic, alpha polypeptide 9	NM_017581	1.43
growth arrest-specific 7	NM_003644	1.43
adaptor-related protein complex 3, sigma 1 subunit	NM_001284	1.43
guanylate cyclase activator 1B (retina)	NM_002098	1.43
Consensus includes gb:AL034396 /DEF=Human DNA sequence from clone 1158B12 on chromosome Xp11.21-11.4 Contains the ZXDA gene for X-linked duplicated Zinc finger A, and MYCL1	AL034396	1.43
Consensus includes gb:AK024584.1 /DEF=Homo sapiens cDNA: FLJ20931 fis, clone ADSE01282. /FEA=mRNA /DB_XREF=gi:10436896 /UG=Hs.203076 Homo sapiens cDNA: FLJ20931 fis,	AK024584	1.43
hypothetical protein DKFZp586M1120	NM_031294	1.43
galanin receptor 1	NM_001480	1.43
melanoma antigen, family B, 2	NM_002364	1.43
KIAA0603 gene product	AJ650848	1.43
Eag-related gene member 2	BC006334	1.43
Consensus includes gb:AA885297 /FEA=EST /DB_XREF=gi:2994374 /DB_XREF=est:aj58h03.s1 /CLONE=IMAGE:1461557 /UG=Hs.323567 CD36 antigen (collagen type I receptor, thrombo	AA885297	1.43
hypothetical protein FLJ10648	NM_018167	1.43
fms-related tyrosine kinase 1 (vascular endothelial growth factor/vascular permeability factor receptor)	NM_002019	1.43
coronin, actin binding protein, 2B	BF939649	1.43
inhibin, beta A (activin A, activin AB alpha polypeptide)	NM_002192	1.43
epithelial protein up-regulated in carcinoma, membrane associated protein 17	NM_005764	1.43
KIAA0241 protein	BE503381	1.43
mitogen-activated protein kinase kinase 7	AF006689	1.43
tight junction protein 1 (zona occludens 1)	NM_003257	1.43
Consensus includes gb:AI050036 /FEA=EST /DB_XREF=gi:3299153 /DB_XREF=est:an31a09.x1 /CLONE=IMAGE:1700248 /UG=Hs.190616 ESTs	AI050036	1.43
cyclin M2	AF216962	1.43
hypothetical protein FLJ14009	AC007766	1.43
protein phosphatase 1, regulatory (inhibitor) subunit 12A	AJ817061	1.43
Consensus includes gb:BF512531 /FEA=EST /DB_XREF=gi:11597710 /DB_XREF=est:UI-H-BW1-amf-b-05-0-UI.s1 /CLONE=IMAGE:3069633 /UG=Hs.297765 ESTs	BF512531	1.43
hairy and enhancer of split (Drosophila) homolog 2	NM_019089	1.43
hypothetical protein FLJ22690	NM_024711	1.43
KIAA1055 protein	AL137303	1.43
MD-2 protein	NM_015364	1.43
core1 UDP-galactose:N-acetylgalactosamine-alpha-R beta 1,3-galactosyltransferase	NM_020156	1.43
arylsulfatase F	NM_004042	1.43
B/K protein	NM_016524	1.43
hypothetical protein MGC3129 similar to ganglioside-induced differentiation-associated protein	NM_024034	1.43
carbonic anhydrase VIII	NM_004056	1.43
allograft inflammatory factor 1	U19713	1.43
hypothetical protein FLJ13954	NM_025020	1.43
Consensus includes gb:AF090101.1 /DEF=Homo sapiens clone IMAGE 30714. /FEA=mRNA /DB_XREF=gi:4063636 /UG=Hs.177837 Homo sapiens clone IMAGE 30714	AF090101	1.43
gb:AF234262.1 /DEF=Homo sapiens IL-1beta-regulated neutrophil survival protein mRNA, complete cds. /FEA=mRNA /PROD=IL-1beta-regulated neutrophil survival protein /DB_XREF=gi:1	AF234262	1.43
hypothetical protein FLJ13842	NM_024645	1.43
syntrophin, gamma 2	NM_018968	1.43
DEAD-box protein	NM_018665	1.43
Consensus includes gb:AJ275374 /DEF=Homo sapiens partial IGVH3 gene for immunoglobulin heavy chain V region, case 1, clone 19 /FEA=CDS /DB_XREF=gi:7573008 /UG=Hs.272357 Ho	AJ275374	1.43
chromosome 8 open reading frame 1	BF057458	1.43

ADP-ribosylarginine hydrolase	NM_001125	1.43
ATPase, Na+/K+ transporting, beta 2 polypeptide	NM_001678	1.43
polyadenylate binding protein-interacting protein 1	AL136920	1.43
hypothetical protein PRO1483	NM_018582	1.43
ERGL protein; ERGIC-53-like protein	NM_021819	1.43
asialoglycoprotein receptor 1	NM_001671	1.43
Consensus includes gb:AK025038.1 /DEF=Homo sapiens cDNA: FLJ21385 fis, clone COL03356. /FEA=mRNA /DB_XREF=gi:10437474 /UG=Hs.306750 Homo sapiens cDNA: FLJ21385 fis, cl	AK025038	1.43
KIAA0530 protein	BF223237	1.43
calcium binding protein 1 (calbrain)	NM_004276	1.43
killer cell immunoglobulin-like receptor, two domains, long cytoplasmic tail, 1	U24078	1.43
myosin VIIA (Usher syndrome 1B (autosomal recessive, severe))	U55209	1.43
integrin, beta 3 (platelet glycoprotein IIIa, antigen CD61)	AI189839	1.43
homeo box D4	NM_014621	1.43
Consensus includes gb:L13283.1 /DEF=Homo sapiens (clone MG2-5-12) mucin (MG2) mRNA, complete polyA site. /FEA=mRNA /DB_XREF=gi:292518 /UG=Hs.103944 Homo sapiens (clon	L13283	1.43
protein kinase, cAMP-dependent, regulatory, type I, beta	AU148827	1.43
ribonuclease, RNase A family, k6	NM_005615	1.43
stem cell growth factor; lymphocyte secreted C-type lectin	D86586	1.43
BMB8 antigen	NM_016564	1.43
RP42 homolog	NM_020640	1.43
Consensus includes gb:AK021842.1 /DEF=Homo sapiens cDNA FLJ11780 fis, clone HEMBA1005931, weakly similar to ZINC FINGER PROTEIN 83. /FEA=mRNA /DB_XREF=gi:10433110 /UG=	AK021842	1.43
thyroid hormone receptor, alpha (erythroblastic leukemia viral (v-erb-a) oncogene homolog, avian)	M24899	1.43
DKFZP586I111 protein	AL050131	1.43
regulator of G-protein signalling 4	BC000737	1.43
Williams-Beuren syndrome chromosome region 5	BC006080	1.43
peroxisome proliferative activated receptor, delta	NM_006238	1.43
NIMA (never in mitosis gene a)-related kinase 2	NM_002497	1.43
hypothetical protein MGC5540	NM_024095	1.43
chromosome X open reading frame 1	NM_004709	1.43
adrenergic, alpha-1A-, receptor	U02569	1.43
glutamate receptor, metabotropic 1	L76631	1.43
death-associated protein kinase 1	NM_004938	1.43
sodium channel, voltage-gated, type II, alpha 2 polypeptide	NM_021007	1.43
putative metallopeptidase (family M19)	NM_022357	1.43
putative non-coding transcript (DiGeorge critical region 5)	AI479306	1.42
nuclear protein, ataxia-telangiectasia locus	U58852	1.42
solute carrier family 5 (neutral amino acid transporters, system A), member 4	AI133127	1.42
likely ortholog of mouse synaptic vesicle glycoprotein 2a	NM_014849	1.42
putative leukocyte platelet-activating factor receptor	M76676	1.42
SNT3 suppressor of mif two 3 homolog 2 (yeast)	AK024823	1.42
corticotropin releasing hormone receptor 1	X72304	1.42
polymerase I and transcript release factor	BC004295	1.42
adenosine A2b receptor	NM_000676	1.42
DHHC1 protein	W61005	1.42
proteasome (prosome, macropain) 26S subunit, non-ATPase, 12	NM_002816	1.42
Consensus includes gb:AC004862 /DEF=Homo sapiens PAC clone RP4-697H17 from 7q11.23-q21.1 /FEA=CDS /DB_XREF=gi:3900856 /UG=Hs.247768 Homo sapiens PAC clone RP4-697H	AC004862	1.42
adenylate cyclase 7	NM_001114	1.42
heterogeneous nuclear ribonucleoprotein L	X16135	1.42
Jagged 1 (Alagille syndrome)	U61276	1.42
Consensus includes gb:AL050047.1 /DEF=Homo sapiens mRNA; cDNA DKFZp5668193 (from clone DKFZp5668193). /FEA=mRNA /DB_XREF=gi:4884286 /UG=Hs.198552 Homo sapiens mR	AL050047	1.42
Consensus includes gb:BE965029 /FEA=EST /DB_XREF=gi:11768976 /DB_XREF=est:601658812R1 /CLONE=IMAGE:3886131 /UG=Hs.198793 Homo sapiens cDNA: FLJ22463 fis, clone HR	BE965029	1.42
Dnal (Hsp40) homolog, subfamily C, member 4	NM_005528	1.42
Consensus includes gb:AK027193.1 /DEF=Homo sapiens cDNA: FLJ23540 fis, clone LNG08239. /FEA=mRNA /DB_XREF=gi:10440262 /UG=Hs.293931 Homo sapiens cDNA: FLJ23540 fis, cl	AK027193	1.42
intersectin 1 (SH3 domain protein)	AF064243	1.42
gonadotropin-releasing hormone receptor	L07949	1.42
KIAA1219 protein	BG236163	1.42
FGFR1 oncogene partner	NM_007045	1.42
C-terminal binding protein 2	N23018	1.42
KIAA1218 protein	AI017382	1.42
solute carrier family 14 (urea transporter), member 2	NM_007163	1.42
fibronectin 1	X02761	1.42
CD163 antigen	Z22970	1.42
myeloproliferative leukemia virus oncogene	NM_005373	1.42
major histocompatibility complex, class II, DO alpha	M29335	1.42

oxidative stress induced like	U46752	1.42
ecotropic viral integration site 2A	NM_014210	1.42
RNA binding motif protein, Y chromosome, family 1, member A1	NM_005058	1.42
src homology three (SH3) and cysteine rich domain	NM_003149	1.42
plexin A2	AI688418	1.42
lipoprotein lipase	BF672975	1.42
homeo box A11	NM_005523	1.42
dihydropyrimidinase-like 3	NM_001387	1.42
hypothetical protein from clones 23549 and 23762	NM_021226	1.42
catenin (cadherin-associated protein), delta 1	AW073672	1.42
ATPase, Ca++ transporting, plasma membrane 3	AW615612	1.42
hypothetical protein FLJ10849	AI539361	1.42
Consensus includes gb:R37655 /FEA=EST /DB_XREF=gi:795111 /DB_XREF=est:yf61d03.s1 /CLONE=IMAGE:26881 /UG=Hs.194761 H.sapiens IMAGE cDNA clone 26881	R37655	1.42
golgi autoantigen, golgin subfamily a, 2	L06147	1.42
hypothetical protein MGC10327	T54159	1.42
interleukin 23, alpha subunit p19	NM_016584	1.42
hypothetical protein FLJ13078	AK023140	1.42
glypican 4	NM_001448	1.42
collagen, type IX, alpha 3	NM_001853	1.42
homeo box (expressed in ES cells) 1	U82811	1.42
cell death-inducing DFFA-like effector b	NM_014430	1.42
KIAA1001 protein	AW052084	1.42
collapsin response mediator protein 1	NM_001313	1.42
potassium inwardly-rectifying channel, subfamily J, member 1	U03884	1.42
hydroxysteroid (17-beta) dehydrogenase 3	NM_000197	1.42
chromosome 21 open reading frame 5	NM_005128	1.42
chromosome 21 open reading frame 2	AI539305	1.42
Consensus includes gb:AA310709 /FEA=EST /DB_XREF=gi:1963058 /DB_XREF=est:EST181521 /UG=Hs.249187 Homo sapiens BAC clone RP11-121A8 from 7p14-p13	AA310709	1.42
attractin	AF106861	1.42
double homeobox, 1	NM_012146	1.42
zinc finger protein 8 (clone HF.18)	AI671579	1.42
neuro-oncological ventral antigen 1	NM_002515	1.42
tumor protein p53-binding protein	AI150117	1.42
transmembrane 4 superfamily member 1	AI189753	1.42
bromodomain-containing 2	S78771	1.42
hypothetical protein FLJ20212	NM_017714	1.42
Consensus includes gb:AK000645.1 /DEF=Homo sapiens cDNA FLJ20638 fis, clone KAT02982, highly similar to SUIS_RABIT SUCRASE-ISOMALTASE. /FEA=mRNA /DB_XREF=gi:7020872 /UG	AK000645	1.42
small inducible cytokine subfamily A (Cys-Cys), member 23	U58913	1.42
Consensus includes gb:BC002889.1 /DEF=Homo sapiens, clone IMAGE:3941350, mRNA, partial cds. /FEA=mRNA /PROD=Unknown (protein for IMAGE:3941350) /DB_XREF=gi:12804072 /L	BC002889	1.42
exostoses (multiple)-like 3	BC006363	1.42
leukocyte immunoglobulin-like receptor, subfamily B (with TM and ITIM domains), member 1	AF009007	1.42
olfactory receptor, family 12, subfamily D member 3	NM_030959	1.42
KIAA0934 protein	N22859	1.42
lymphocyte antigen 6 complex, locus E	NM_002346	1.42
Consensus includes gb:AI983505 /FEA=EST /DB_XREF=gi:5810724 /DB_XREF=est:wt49h12.x1 /CLONE=IMAGE:2510855 /UG=Hs.137576 ribosomal protein L34 pseudogene 1	AI983505	1.42
phosphodiesterase 4C, cAMP-specific (phosphodiesterase E1 dunce homolog, Drosophila)	U88712	1.42
CD44 antigen (homing function and Indian blood group system)	M24915	1.42
hypothetical protein FLJ10512	AI655902	1.42
wingless-type MMTV integration site family, member 4	NM_030761	1.42
galactosamine (N-acetyl)-6-sulfate sulfatase (Morquio syndrome, mucopolysaccharidosis type IVA)	NM_000512	1.42
MCF.2 cell line derived transforming sequence-like	AB002360	1.42
arginine vasopressin receptor 2 (nephrogenic diabetes insipidus)	NM_000054	1.42
mitochondrial solute carrier	NM_018579	1.42
cytotoxic T-lymphocyte-associated protein 4	NM_005214	1.42
glutamine-fructose-6-phosphate transaminase 2	NM_005110	1.42
Consensus includes gb:AI275355 /DEF=Homo sapiens partial IGVH3 V3-20 gene for immunoglobulin heavy chain V region, case 1, clone 2 /FEA=CDS /DB_XREF=gi:7572968 /UG=Hs.2723	AI275355	1.42
KIAA0459 protein	AL537463	1.42
tumor necrosis factor receptor superfamily, member 10d, decoy with truncated death domain	AF021233	1.42
zinc finger protein 272	X78931	1.42
collagen, type V, alpha 1	AI862325	1.42
hypothetical protein FLJ20220	NM_017718	1.42
parathyroid hormone-like hormone	NM_002820	1.42
solute carrier family 24 (sodium/potassium/calcium exchanger), member 1	AF026132	1.42

cytochrome b-561	NM_001915	1.42
ring finger protein (C3HC4 type) 8	NM_003958	1.42
SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily b, member 1	NM_003073	1.42
ATPase, Ca++ transporting, cardiac muscle, fast twitch 1	NM_004320	1.42
H2B histone family, member J	NM_003524	1.42
KIAA1193 protein	AB033019	1.42
5-hydroxytryptamine (serotonin) receptor 1F	NM_000866	1.42
tetraspan transmembrane 4 super family	NM_006675	1.42
GTP-binding protein	U69563	1.42
Consensus includes gb:AV720803 /FEA=EST /DB_XREF=gi:10817955 /DB_XREF=est:AV720803 /CLONE=GLCCMA04 /UG=Hs.293055 ESTs	AV720803	1.42
neutrophil cytosolic factor 2 (65kD, chronic granulomatous disease, autosomal 2)	BC001606	1.42
ladinin 1	NM_005558	1.42
KIAA0643 protein	NM_024793	1.42
hypothetical protein MGC4638	BC005161	1.42
ribosomal protein S6 kinase, 90kD, polypeptide 6	NM_014496	1.42
Consensus includes gb:AI307760 /FEA=EST /DB_XREF=gi:4002364 /DB_XREF=est:tb24g09.x1 /CLONE=IMAGE:2055328 /UG=Hs.14074 Homo sapiens clone 24627 mRNA sequence	AI307760	1.42
pregnancy-induced growth inhibitor	NM_013370	1.42
KIAA0830 protein	AL573201	1.42
Consensus includes gb:AL049988.1 /DEF=Homo sapiens mRNA; cDNA DKFZp564F212 (from clone DKFZp564F212). /FEA=mRNA /DB_XREF=gi:4884239 /UG=Hs.306304 Homo sapiens mF	AL049988	1.42
hypothetical protein FLJ10986	NM_018291	1.42
DKFZP434M154 protein	AI401612	1.42
TGFB inducible early growth response 2	AA149594	1.42
RAB2, member RAS oncogene family-like	AL050259	1.42
cytochrome P450, subfamily IIA (phenobarbital-inducible), polypeptide 6	NM_000762	1.42
hypothetical protein	NM_019073	1.42
matrix metalloproteinase 16 (membrane-inserted)	AB009303	1.42
Consensus includes gb:AW517464 /FEA=EST /DB_XREF=gi:7155546 /DB_XREF=est:xp94g06.x1 /CLONE=IMAGE:2748058 /UG=Hs.119598 ribosomal protein L3	AW517464	1.42
cannabinoid receptor 2 (macrophage)	NM_001841	1.42
cytokeratin 20	AI732381	1.42
thioredoxin-like 2	NM_006541	1.42
Consensus includes gb:AL080072.1 /DEF=Homo sapiens mRNA; cDNA DKFZp564M0616 (from clone DKFZp564M0616). /FEA=mRNA /DB_XREF=gi:5262482 /UG=Hs.21195 Homo sapiens ı	AL080072	1.42
membrane protein, palmitoylated 2 (MAGUK p55 subfamily member 2)	AW450911	1.42
interleukin 10	NM_000572	1.42
adenylate kinase 1	BC001116	1.42
MUF1 protein	AK024051	1.42
myogenic factor 3	NM_002478	1.42
carboxypeptidase E	NM_001873	1.42
KIAA0523 protein	NM_014021	1.42
Kell blood group	NM_000420	1.42
Ras homolog enriched in brain 2	D78132	1.42
FK506 binding protein 9 (63 kD)	AL050187	1.42
A kinase (PRKA) anchor protein 7	AF161075	1.42
KIAA1052 protein	AI655714	1.42
adenosine A1 receptor	NM_000674	1.42
CCR4-NOT transcription complex, subunit 2	AI123426	1.42
Werner helicase interacting protein	NM_020135	1.42
hypothetical protein FLJ13511	R77097	1.42
hypothetical protein PRO2325	NM_018523	1.42
KIAA0410 gene product	NM_014778	1.42
plexin B2	AF336795	1.42
GATA binding protein 3	BC003070	1.42
M-phase phosphoprotein 1	NM_016195	1.42
a disintegrin-like and metalloprotease (reprolysin type) with thrombospondin type 1 motif, 9	NM_020249	1.42
calcium channel, voltage-dependent, L type, alpha 1B subunit	NM_000718	1.42
epilepsy, progressive myoclonus type 2, Lafora disease (laforin)	AK022721	1.42
methionine adenosyltransferase II, beta	NM_013283	1.42
Consensus includes gb:AL137606.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434H0211 (from clone DKFZp434H0211). /FEA=mRNA /DB_XREF=gi:6808354 /UG=Hs.151171 Homo sapiens	AL137606	1.42
metastatin 1	N32151	1.42
Consensus includes gb:BG545769 /FEA=EST /DB_XREF=gi:13544434 /DB_XREF=est:602573042F1 /CLONE=IMAGE:4701250 /UG=Hs.9927 Homo sapiens mRNA; cDNA DKFZp564D156 (fr	BG545769	1.42
growth arrest-specific 7	NM_005890	1.42
Consensus includes gb:AK025363.1 /DEF=Homo sapiens cDNA: FLJ21710 fis, clone COL10087. /FEA=mRNA /DB_XREF=gi:10437865 /UG=Hs.306807 Homo sapiens cDNA: FLJ21710 fis, cl	AK025363	1.42
enolase alpha, lung-specific	NM_014355	1.42
calcium/calmodulin-dependent protein kinase (CaM kinase) II beta	AF078803	1.42

adrenomedullin	NM_001124	1.42
PRO0628 protein	NM_014134	1.42
transcobalamin II; macrocytic anemia	NM_000355	1.42
Consensus includes gb:AI879383 /FEA=EST /DB_XREF=gi:5553432 /DB_XREF=est:au50b05.x1 /CLONE=IMAGE:2518161 /UG=Hs.190885 ESTs, Moderately similar to ALU1_HUMAN ALU SI	AI879383	1.42
myosin VA (heavy polypeptide 12, myoxin)	Z22957	1.42
Consensus includes gb:BF574664 /FEA=EST /DB_XREF=gi:11648376 /DB_XREF=est:602132139F1 /CLONE=IMAGE:4271792 /UG=Hs.274531 Homo sapiens mRNA; cDNA DKFZp586F1024	BF574664	1.42
Consensus includes gb:AA654415 /FEA=EST /DB_XREF=gi:2590569 /DB_XREF=est:nt03g09.s1 /CLONE=IMAGE:1192096 /UG=Hs.322808 ESTs, Moderately similar to AMYP_HUMAN ALPHA	AA654415	1.42
lymphoid blast crisis oncogene	AF126008	1.42
chromosome 1 open reading frame 17	AF288389	1.42
melanoma antigen, family A, 5	AI200443	1.42
Consensus includes gb:AK000257.1 /DEF=Homo sapiens cDNA FLJ20250 fis, clone COLF6635. /FEA=mRNA /DB_XREF=gi:7020217 /UG=Hs.274505 Homo sapiens mRNA; cDNA DKFZp564A	AK000257	1.42
UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase 3 (GalNAc-T3)	BF063271	1.42
PTK2 protein tyrosine kinase 2	NM_005607	1.42
Friend of GATA2	NM_012082	1.42
tumor necrosis factor receptor superfamily, member 6b, decoy	NM_003823	1.42
BUB1 budding uninhibited by benzimidazoles 1 homolog (yeast)	AU156551	1.42
discoidin domain receptor family, member 1	NM_001954	1.42
hypothetical protein FLJ23033	NM_024686	1.42
hypothetical protein FLJ10063	NM_017987	1.42
Consensus includes gb:BG111761 /FEA=EST /DB_XREF=gi:12605267 /DB_XREF=est:602285343F1 /CLONE=IMAGE:4372619 /UG=Hs.8107 Homo sapiens mRNA; cDNA DKFZp58680918 (I	BG111761	1.42
Consensus includes gb:AI273812 /FEA=EST /DB_XREF=gi:3896080 /DB_XREF=est:qu08t07.x1 /CLONE=IMAGE:1964197 /UG=Hs.169252 Human L-myc protein gene, complete cds	AI273812	1.41
Consensus includes gb:AL110227.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434J194 (from clone DKFZp434J194). /FEA=mRNA /DB_XREF=gi:5817165 /UG=Hs.323067 Homo sapiens mRI	AL110227	1.41
hydroxyacyl-Coenzyme A dehydrogenase/3-ketoacyl-Coenzyme A thiolase/enoyl-Coenzyme A hydratase (trifunctional protein), alpha subunit	U04628	1.41
neuroligin 1	NM_014932	1.41
Eag-related gene member 2	NM_030779	1.41
Consensus includes gb:AU145746 /FEA=EST /DB_XREF=gi:11007267 /DB_XREF=est:AU145746 /CLONE=HEMBA1005679 /UG=Hs.82193 esterase D formylglutathione hydrolase	AU145746	1.41
similar to S. pombe dim1+	BC001046	1.41
Consensus includes gb:AF070647.1 /DEF=Homo sapiens clone 24438 mRNA sequence. /FEA=mRNA /DB_XREF=gi:3283921 /UG=Hs.124126 Homo sapiens clone 24438 mRNA sequence	AF070647	1.41
Wiskott-Aldrich syndrome-like	BE504979	1.41
hypothetical protein FLJ14327	NM_024912	1.41
interferon, beta 1, fibroblast	NM_002176	1.41
myotubular myopathy 1	U46024	1.41
retinal short-chain dehydrogenase/reductase retSDR2	NM_016245	1.41
BCL2-related protein A1	NM_004049	1.41
Rho GTPase activating protein 6	NM_001174	1.41
solute carrier family 17 (anion/sugar transporter), member 5	NM_012434	1.41
ets variant gene 6 (TEL oncogene)	AF041811	1.41
H2B histone family, member D	NM_003520	1.41
radial spokehead-like 1	NM_030785	1.41
Consensus includes gb:BE930512 /FEA=EST /DB_XREF=gi:10456588 /DB_XREF=est:RC6-GN0071-160800-021-D01 /UG=Hs.168732 ESTs	BE930512	1.41
lethal (3) malignant brain tumor l(3)mbt protein (Drosophila) homolog	AW445040	1.41
hypothetical protein FLJ10698	AI951798	1.41
arylacetamide deacetylase (esterase)	NM_001086	1.41
Consensus includes gb:AK024108.1 /DEF=Homo sapiens cDNA FLJ14046 fis, clone HEMBA1006461. /FEA=mRNA /DB_XREF=gi:10436406 /UG=Hs.142677 Homo sapiens cDNA FLJ14046 fi	AK024108	1.41
transducin-like enhancer of split 2 (E(sp1) homolog, Drosophila)	M99436	1.41
hypothetical protein MGC3136	NM_024105	1.41
nuclear RNA export factor 3	NM_022052	1.41
basic-helix-loop-helix-PAS protein	NM_022123	1.41
Consensus includes gb:Z34282.1 /DEF=H.sapiens (MAR11) MUC5AC mRNA for mucin (partial). /FEA=mRNA /GEN=MUC5AC /PROD=mucin /DB_XREF=gi:563384 /UG=Hs.247953 H.sapiens	Z34282	1.41
Consensus includes gb:BF223401 /FEA=EST /DB_XREF=gi:11130578 /DB_XREF=est:7q87h04.x1 /CLONE=IMAGE:3705462 /UG=Hs.112158 Homo sapiens PAC clone RP4-751H13 from 7q	BF223401	1.41
Consensus includes gb:U63332.1 /DEF=Human super cysteine rich protein mRNA, partial cds. /FEA=mRNA /PROD=super cysteine rich protein /DB_XREF=gi:1480862 /UG=Hs.169325 Hum	U63332	1.41
KIAA0750 gene product	NM_014632	1.41
KIAA0241 protein	AI797458	1.41
ribotinase	AI823980	1.41
Consensus includes gb:AL050376.1 /DEF=Homo sapiens mRNA; cDNA DKFZp586J101 (from clone DKFZp586J101). /FEA=mRNA /DB_XREF=gi:4914609 /UG=Hs.322645 Homo sapiens mRI	AL050376	1.41
hypothetical protein FLJ11703	NM_024969	1.41
tolloid-like 2	AA582404	1.41
biliverdin reductase A	U34877	1.41
hypothetical protein IMAGE:4215339	AV721013	1.41
kinesin-like 3	NM_030615	1.41
BTG3 associated nuclear protein	AK023827	1.41
parvalbumin	NM_002854	1.41
opiate receptor-like 1	NM_000913	1.41